

AGTCCGGGGG	ACCTTTTTAG	TCGGTAGATT	GAGATTGCAA	ACAAATCTGC	ATCTACATTG	4860
AAAGCTTAAT	TTCTAATAAT	TGAAAAAATC	GAATGAAAAA	TTTCTTACCT	TCATTACACAG	4920
AGCTCGATTT	CAGAGCTCTT	TTTGCTAGCT	TATTCATACT	TTTCTGAATT	TCGAAAAAGA	4980
AATGTAAGCG	TTTGATAGAT	TTACAAAAAG	ATTGTATAAT	AGGGATAAGA	ATAGAAAAGG	5040
AGAAGTCTCA	TGGCAGTTAA	AGATTTTATG	ACCCGCAAGG	TAGTTTATAT	TAGTCCAGAT	5100
ATAACAGTAT	CTCATGCAGC	AGATTTGATG	AGAGAGCAAG	GTTTGCACCG	TCTGCCTGTT	5160
ATCGAAAATG	ATCAATTAGT	TGGTTTGGTG	ACTGAGGGAA	CCATTGCACA	AGCAAGTCCA	5220
TCTAAAGCAA	CAAGTCTTTC	TATCTATGAG	ATGAATTATC	TTCTGAATAA	GACAAAAGTA	5280
AAAGATGTCA	TGATTTCGCGA	TGTTGTCACT	GTCTCAGGCT	ATGCTAGTCT	AGAAGATGCA	5340
ACTTATCTGA	TGTTGAAAAA	TAAGATTAGT	ATTCTCCCTG	TCGTAGATAA	CCATCAAGTA	5400
TACGGAGTTA	TTACTGACCG	TGACGTTTTC	CAAGCCTTTC	TTGAAATTGC	AGGTTATGGC	5460
GAAGAAGGGA	TTCTGTACG	CTTTGTTACA	GAAGATGAAG	TTGGTGTTC	TGAAAAAATT	5520
GTTTCTTTGA	TTGTAGAAGA	AAATTTGAAT	ATCTCCATA	CAGTCAATAT	TCCGCGTAAG	5580
GATGGTAAGG	TGATTATCGA	AGTGCAAATC	GATGGATCAA	TTGATTTACC	AGCCTTGAAA	5640
GAAAAATTTG	AAGCAAATGG	TATTCAAGTG	GAAGAAATCG	CTCGCACTTC	AGCAAAAGTC	5700
TTGTAAAGAAG	GGAAGCCCAA	AGGCTTCTTT	TTTCATGAAA	AGGGGATTAG	AGCAAAAGAT	5760
GGAAAGAAAT	GATAAAATAT	GCTATAATGA	AATAATGTAA	AAAAGGAGTA	TTTATGGACA	5820
TTTCAGTAAT	TCGTACAGAA	ATTGACGCAA	ATCGTGAAAA	ATTAGCTTCT	TTCAGGGGGT	5880
CTCTTTGACC	TCGAAGGGCT	AGAGGAAGAG	ATTGCCATCT	TGAAAAACAA	GATGACAGAA	5940
CCTGATTTTT	GGAACGATAA	TATTGCGGCC	CAAAAACGT	CGCAAGAATT	AAATGAATTA	6000
AAAAACACTT	ACAATACCTT	CCATAAGATG	GAAGAGTTGC	AGGATGAAGT	CGAAATTTTA	6060
TTGGATTTTT	TGGCTGAAGA	CGAGTCAGTG	CATGATGAAC	TGGTAGCGCA	GTTAGCCGAA	6120
CTTGATAAGA	TAATGACCAG	CTACGAGATG	ACTCTACTCT	TGTCAGAACC	TTATGACCAC	6180
AACAATGCCA	TCTTGGAAT	CCATCCAGGT	TCTGGTGGTA	CTGAGGCGCA	GGACTGGGGT	6240
GATATGTTGC	TTCTGATGTA	TACTCGTTAT	GGTAATGCTA	AAGGCTTTAA	AGTGGAAGTG	6300
TTGGATTACC	AAGCAGGTGA	TGAGGCTGGT	ATTAAGTCGG	TAACTTTATC	ATTTGAAGGG	6360
CCTAATGCCT	ATGGTCTCCT	CAAGTCAGAA	ATGGGTGTTT	ACCGCTTAGT	GCGAATCTCA	6420
CCATTTGACT	CTGCCAAACG	TCGCCATACC	TCTTTCACAT	CTGTAGAAGT	GATGCCAGAA	6480
TTGGATGATA	CTATPGAAGT	GGAAATCCGT	GAAGATGATA	TCAAGATGGA	TACCTTCCGT	6540

750

TCAGGTGGTG CCGGTGGACA AAACGTCAAT AAGGTTTCAA CAGGTGTACG TTTAACCCAC 6600
 ATTCCAACCTG GAATTGTTGT CCAATCAACA GTAGATCGTA CCCAGTATGG AAATAGAGAT 6660
 CGTGCCATGA AGATGTTGCA GGCTAAGCTC TATCAAATGG AGCAAGATAA GAAGGCTGCG 6720
 GAGGTAGATT CTCTCAAAGG TGAGAAAAAG GAGATCACTT GGGGAAGCCA AATCCGTTCT 6780
 TATGTCTTCA CGCCTTATAC TATGGTAAAA GATCACCGAA CTAGCTTTGA GGTGCTCAG 6840
 GTAGATAAGG TTATGGATGG GGACCTAGAT GGTTTTATCG ATGCTTATCT CAAGTGGCGA 6900
 ATTAGCTAAG ATAGAAAAGG ACTCACATGT CAATTATTGA AATGAGAGAT GTCGTTAAAA 6960
 AATACGACAA CGGAACAACCT GCTCTACGCG GTGTTTCGGT TAGCGTTCAA CCGGGGGAAT 7020
 TTGCTTACAT CGTAGGACCT TCAGGAGCAG GGAAGTCAAC TTTTATTCGT TCTCTGTATC 7080
 GTGAAGTAAA AATCGATAAA GGAAGCCTAT CAGTTGCTGG TTTTAATCTG GTTAAGATCA 7140
 AAAAGAAAGA TGTCCCGCTT CTACGTCGTA GTGTTGGGGT TGTCTTCCAG GATTATAAAT 7200
 TGTTACCAA GAAAACCTGTC TATGAAAATA TTGCTTACGC TATGGAAAGTA ATCGGGGAAA 7260
 ATCGCCGTAA TATCAAAGA CGAGTGATGG AAGTTTGGGA CTTGGTTGGA TTGAAGCATA 7320
 AGGTTTCGTT TTTCCCAAT GAACCTCAG GTGGGGAGCA ACAGCGGATT GCGATTGCGC 7380
 GTGCAATTGT AAATAATCCC AAAGTATTGA TAGCTGATGA GCCAACAGGA AATCTGGATC 7440
 CGGATAATTC ATGGGAAAT ATGAATCTCT TGGAACGGAT TAACyTACAA GGAACAACCTA 7500
 TTTTGATGGC GACTCATAAT AGCCAGATTG TAAATACCTT GCGCCACCGT GTCATTGCCA 7560
 TTGAAAATGG CCGTGTGCTT CGTGACGAAT CAAAAGGAGA GTATGGATAC GATGATTAGT 7620
 AGATTTTTTC GCCATTTAT TGAAGCCTTA AAAAGTTTGA AACGAAATGG TTGGATGACA 7680
 GTAGCTGCTG TCAGTTCAGT CATGATTACT TTGACCTTGG TGGCAATAT TGCATCTGTT 7740
 ATTTTCAATA CAGCGAAACT AGCTACAGAT ATTGAAAATA ATGTCCGTGT AGTAGTTTAT 7800
 ATCCGAAAGG ATGTGGAAGA TAATAGTCAG ACAATTGAAA AAGAAGGTCA AACTGTTACA 7860
 AATAATGACT ACCACAAGGT ATATGATTCT TTGAAGAACA TGTCTACGGT TAAAAGTGTT 7920
 ACCTTTTCAA GTAAAGAAGA ACAATATGAA AAATTAACCG AGATAATGGG AGATAACTGG 7980
 AAAATCTTTG AAGGAGATGC CAATCCTCTC TATGATGCCT ATATTGTAGA GGCAAACACT 8040
 CCAAATGATG TAAAAACTAT AGCCGAAGAT GCTAAAAAAA TTGAAGGTGT CTCTGAGGTT 8100
 CAAGATGGCG GTGCCAATAC AGAAAGACTC TTCAAGTTAG CTTCATTTAT CCGTGTTTGG 8160
 GGACTAGGGA TTGCTGCTTT GTTAATTTTT ATCGCAGTTT TCTTGATTTT AAATACCATT 8220
 CGTATTACCA TTATTTCCCG CAGTCGCGAA ATTCAAATCA TCGCCTTGGT CCGAGCTAAA 8280
 AACAGTTATA TCCGTGGACC GTTCTTGTTA GAAGGAGCCT TTATCGGTTT ATTGGGAGCT 8340

751

ATCGCACCAT CTGTTTGGT CTTTATTGTT TATCAAATTG TTTACCAATC TGCAACAAA	8400
TCGTTGGTAG GGC AAAATCT ATCCATGATT AGTCCAGATT TATTTAGTCC GTTGATGATT	8460
GCCCTACTAT TTGTGATTGG GGTTCATT GGTTCATTGG GATCAGGAAT ATCCATGCGC	8520
CGATTCTTGA AGATTAGGT AAAATAGCTG CTTTTATGAG GAGATTGTAA AATCTCCTTT	8580
TTTGCTACAA GAGTTTTGA AAAGAGATGC GCAGAAGAAA AGAGCTTCCA AAGAAGTCCC	8640
CCAGAGAAGA CTTC	8654

(2) INFORMATION FOR SEQ ID NO: 99:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 19718 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 99:

TGTCGCGTCA AAATCATTAC TATGGCTATG TATAGCCCTT ACTATGACTT GGCTAAACAC	60
GTTCGCTTTC AAATTTCTAG GCTCAGGCTG AACAGTCTC CCAGGCTGTT CACTCCCGAA	120
TGCTAAAATC GTTCTTGATC GCTTTCACAT TGTACAACAT CTTAGCCGTG CTATGAGTCG	180
TGTGCATGTC CAAATCATGA ATCAGTTTCA TCGAAAATCC CATGAATACA AGGCTATCAA	240
GCGCTACTGG AAACATTC AACAGGATAG CCGTAAACTG AGTGATAAGC GATTTTATCG	300
CCCTACTTTT CGCATGCACT TAACAAATAA AGAAATCTT GACAAGATTT TAAGCTATTC	360
AGAAGACTTG AAACACCACT ATCAGATCTA TCAACTCTTA CTTTTTCACT TTCAGAACAA	420
AGACCCTGAG AAATTTTTCG GACTCATTGA GGACAATCTG AAGCAGGTTT ATCCTCTTTT	480
TCAGACTGTC TTTAAAACCT TTCTCAAAGA TAAAGAAAAG ATTATCAACG CCCTTCAACT	540
ACACTATTCT AATGCCAAAC TGAAGCGAC CAATAATCTC ATCAAACCTA TCAAGCGCAA	600
TGCCTTTGGT TTTTCGAACT TTGAAAACCT CAAAAACGG ATTTTATCG CTTTGAACAT	660
CAAAAAGAA AGGACGAAAT TTGTCCTTTC TCGAGCTTAG CTGACTTCAA CCCACTACAG	720
TTGACAAAGA GCCTAATTTT CATAAAAATT GACATGGAAA TTATAAAACC ATTACTAGTT	780
TAGTCTTTT TGATAACGTG CCAATTCGGC TTGGTTTCGCC CAAACATAGT GACCTGGACG	840
GATTTCTACC ATAGATGGCT TATCAGTCTC ATAGTCGTGT TGAAGTGGAT CGTAAACCTT	900
CAAGACCTTC TTACGTTCCA AGATTGGATC TGGGATTGGT ACCGCTGAAA GCAAGGCTTG	960
AGTATATGGG TGAATGGAT TGTAAACAA TTCTTCTGTT TCTGCAACCT CTACAATAAC	1020

752

ACCCTTGTA	ATAACTGCGA	TACGATCTGA	AATAAAGCGA	ACAACCGACA	AGTCATGGGC	1080
GATGAAGAGA	TAGGTCAGGC	CGAGCTCTTT	TTGGAATTTT	TTGAGCAAGT	TCAAGACTTG	1140
GGCACGTACA	GAAACGTCCA	AGGCTGAAAT	TGGCTCATCT	GCAATAACAA	AGTCTGGTTG	1200
CATGACCAAG	GCACGGGCAA	TACCGATACG	TTGACGTTGA	CCGCCTGAGA	ATTCATGAGG	1260
GTAACGAGTC	AAGTGTCTCAG	CAAGAAGACC	TACTTCACGG	ATAATATTTT	GAAC'TTCTC	1320
TTTACGTTCT	TCTTCATCCT	TAAATAAACG	GTGATTGTAA	AGACCTTCAG	AAATAATATA	1380
ATCAACAGTC	GCACGTTTCT	TCAAACCTGC	GGCAGGGTCT	TGGAAAATCA	TCTGGATTCTG	1440
ACGAATCAAT	TCCGCAGCTT	GTTACGCGA	TTTCTTACCA	TTAATCTTTT	GACCATCAAA	1500
AATGATATCT	CCATFACTTG	TATCATTTAG	ACCGATGATA	GCACGACCAA	TAGTTGTTTT	1560
CCCCTACCG	GACTCACCTA	CAAGCGAGAA	AGTTTCTCCC	TTGTTGATAA	AGAAGTTAGC	1620
ATTTTTAACC	GCGACAAACT	TCTTACTTCC	TTCACCGAAG	GAAATTTCTA	AATCTTTGAT	1680
TTCTACTAAT	TTTTTCAGACA	TTTCCTTCCT	CCTAGTCAGC	CAGATGGGCA	AATCCCATTT	1740
TTTCACGGAT	CTTATCATGG	AGATTGCAA	TCACAGCTGG	TTTTTCTACT	TTCGGAGCAT	1800
CCTCATGAAG	AAGCCAAGTT	TTAGCCCAAT	GTGTCTCTGA	TACTGAGAAT	TGAGGAGCTT	1860
TTTGTTGAA	GTCAATCTGC	ATTGCGTAGT	CAGAACGCAA	GGCAAAAGCA	TCCCCTTTCA	1920
GGTCAGTATA	AAGTGACGGA	GGTGTTCCTG	GGATTGAGTA	AAGATCCCCCT	TTATCATCAG	1980
CAAGCTGAGG	CAAGCTAGAC	AAGAGACTCC	ATGTATATGG	ATGGCGAGGG	TCATAGAAGA	2040
CTTCTCAAC	CGTCCATAC	TCAACGATTT	CTCCTGCATA	CATAACCGCT	ACCTTATCCG	2100
CAATACTTGC	CACCACACCA	AGGTCGTGGG	TAATAAAGAT	TGTTGTGAAA	TGATACTCGT	2160
TTTGTAAGA	TTTTAGCAAA	TCAATAATCT	GAGCTTGAAT	AGTTACATCC	AAGGCAGTTG	2220
TTGGCTCATC	ACAGATCAAG	ACATCAGGTC	GGCAGGCAAG	GGCAATAGCA	ATAACGATAC	2280
GTGACGCAT	TCCTCCAGAA	TATTGGAATG	GGTATTCATT	AAAACGTCTA	TCTGCGTCTG	2340
GAATGCCAAC	CTTATTCATG	TAGTCAATGG	CCAATTCCTT	CGCTTCTTTA	GCTGTTTTTC	2400
CTTGGTGTTC	TACAATAACT	TCTGTAATCT	GACTACCAAT	TGTTTTAATG	GGGTCCAAAC	2460
TAGTCATTGG	GTCTTGAAG	ATAGTCGCAA	TCTTAGCACC	ACGAATTTGT	TCCAATCCT	2520
TGTGAGAAGA	TAAAGCTGTC	AAGTCCTGAC	CACGGTAGTC	AATACTACCT	TGGGCAATAC	2580
GACCATTTTC	TTCGAGCATA	CCTGTGAAGG	TCTTGTCAA	AACAGATTTA	CCTGATCCTG	2640
ACTCACCTAC	CAAGGCTAAT	ACTTCTCCTT	CGACTAGTTC	AAGGGAAACG	CCGCGAATGG	2700
CTGTCAATAC	TTTGTACGA	ACGTCAAAT	CCACGACAAT	ATCGCGAGCA	GTCAAAATTA	2760
CATTTTTTTC	TTTTGTCAAT	TCTACTCCTA	TCTATGTGTA	CGTGGATCAC	TAGCATCCGC	2820

753

TAAGTTTTGA CCAACTACGA AAAGGGACAA GGATACCAAG ACAAGGGTTG TCAATGGAAT	2880
CCAGAACAAG TAAGCATGG TTGTTACGTT TTGTGAATAA TCCGAAATCA AACGACCCAA	2940
ACTTGGCACT GTAATCGGTA ATCCAAGACC GAAGAAAGAC AAGAAGGCTT CGTATGAGAT	3000
AAAGCTTGGA AGCATTTGAG TCATGGTGT CACAATAACA GATACCAATT GAGGCATGAT	3060
ATTTTTGGCA ACAATCTTCA AGGTTGGTGT TCCCAAAGTA CGTGACGCCA AGTTGTATTC	3120
CAAGTCACGA TAGCGCAAGA TTTGCACACG GATCATGAAG GCAATACCAA TCCATGTTGT	3180
TACGCTCATG GCAAAAATCA GATTCCAGAA TCCAGCTCCG ATTGAGTAAG TCAAGACAAT	3240
AACAATCAAA AGAGGTGGGA TGTTTGAGAT GACGTTGTAA ACTTCCATCA TGACACGGTC	3300
AACTGATTTT GAAATACCCC AAATACCACC GACAAAACA CCGATAACCA AGTTAATCAC	3360
TGTCGCAATC ACAGAAATGA GGATGGAGTT ACGAGCTCCG AACCAGACAC CGTCAAAGAG	3420
CGATTTACCG TTAAGTCTAG TACCGAACCA ATGCTCCGCA TTTGGCTTGA TATAACGAAC	3480
ACTAAAGTCG TTTACCTTGC TGACATCATT GAAATCAAAC TTAGAAAACA TTGGGTAGAT	3540
GAAACTTATC AAAATGATGG CTACCAAGAT TCCCAACATG ACTACAGTTG ATTTTTTCTT	3600
CATAAATGT TTAACACTG ATTTCCAGTA AGAATATGCT GGCGCATCAA TAGTTTCAGA	3660
GGCAAAATCG TCACGTTTAA CAAACTGAAA TTTTCTTTA TCGATTGTAG ACATTATTTG	3720
CCTCCTTCT CAGTCAATTT AATACGTGGG TCAATAATAG TCATCCAAAT ATCTCCCAA	3780
AGACGTGAGA AGATAGAAAT ACATGTAAAG ATGAAGACAA GACCAACGAC CATAGAGTTA	3840
TTAGATGCTT TTACAGAGTC AATCAACATT TTACCCATAC CTGGGAAGGC GAAGACTGTT	3900
TCAGTAAGGG TTGCACCACC GATAACCCCA ATAATGGCAG CAGGAATCC TGAAACCAGC	3960
GGAACCATGG CATTTTAAA GATGTGTTG TTTGAAATTT CTTTTTCAGA CAAACCTTTT	4020
GCACGAGCGA AACGAACAAA GTCTTGAGAT TGCAAGTCAA TCATGTAACG ACGAATCCAA	4080
ATGGCTGTAC CAGGAGCACC CAACAAACCA AGGATGACTG CTGGTAAAAC GTAAGAACGC	4140
CAATCTCCAG CTCCCAAGAT AGGGAATGAA TCTGGAAGGG CAATAGATGA TCCAATCAAT	4200
CGAACGATGT AAACCAAGGC AATCGTTGGA AGAGCAAGCA AGAAGGTCAA AGCCCCGTT	4260
GAGAGGCTAT CAATCCAAGT GTTCTTGAAA CGAGCCATGG CTGAACCAAG TGGCACGGCA	4320
AGAGCATAGG CAAGAACCAA ACCAATCAAA CCAGTAATAG CAGAGCTGAC AATCATAGAT	4380
GGATATTGGT AATTACTTTC AGTCGCTGTA TAAGGATCAT CTTCCATA GCTAGCTACT	4440
TCACGAGAGT CAGCCTGACT AGGTGACTTG TAGGTTCTTG AGTAAATATT TACAGAAGAC	4500
GTTTTCTTAC CTGTTGGGAA CTGAACTTGG GCAGTTTGG TTTGTCCTTG ACCTTGAGTA	4560

754

ATAACCTGAA	GAACCTGGTGT	ATTAGCATAG	GTTGGGTAAG	AGTCACCTAA	ATTCAAGTTC	4620
ACAAAGTTTT	GATGAACAAA	TGGGAAGTGA	CTGTTAAAGT	ACAAGAGATA	TTTATGTTTA	4680
GTTCCCTGAAC	CGACCAATGA	CCATCCGATA	GCTGGATCAT	TTTCAAAAACG	AAGGTAGCGT	4740
TTCAAGTCTG	GATTTTCAGG	GTCTTGGATT	TTATTTGTAT	GGTCAATGTC	AATCAAGTTA	4800
GCATAGAAGT	GAAAAACACG	TTCAAAAATT	GGAATTTTAC	GAGTAGCATA	GAATTGACCA	4860
CTTTCAGTAA	ATTCTCCCAA	AGTCCAACCA	TGACCTAATT	GATTGATGTA	CTTTTCATAA	4920
ATAGCTTTAT	TGGTCGCATT	TGCTTCTACT	GTTACAGAAG	AATCCATGCT	ACTTGCCTTT	4980
TCTTGCAACT	CTTTAGTATC	GTAATACTCA	ATGTAGCCCA	TACGCTCAA	CACAGTATTT	5040
TCATAGTTAT	CACGTTTATC	AGCCGTGTGC	GCAATTTTAT	TATAGTTAGG	ATCCTGCTTG	5100
AAAATCAATT	TTCGAGGAAC	CAAGGTATAG	ATAATCGTGT	AGGTCAAAGT	CGTTACTAAG	5160
AAAATCGAAA	CCAATGACCG	CAAAACACGC	ATAAAAATAT	ATTTTTTCAT	ATTATTTCCCT	5220
TTAAAAATCC	CAAAAGAACC	TTCTCCTCAT	GGAGAGAAA	TTCTATTAGA	AATTATTTAC	5280
TTCACATGAC	TTGCCAATTC	TTTTTGAGCT	TTCTCATTTG	ATTCAGCTTT	TTCTTTCAAC	5340
CATTTTTCAC	GAGCTTTTTC	ATACTCTTCC	TTAGTCACCA	CTTTATCTTG	TGATTTCAA	5400
TATTTGAAGT	AAACATCTGA	CCCCTTAGAG	CCTGTTTGGC	CAGAAGCTCC	AGTAAATGGA	5460
ACAATTCGTG	AAAGCACTGG	TGCTGCACCA	GAAGAAGCCA	TAGCAGGAAT	AAAGAGTGAA	5520
CTATCTGTCA	ACCATGCTTG	AGCCGCTGCA	TATTTTTTCAT	AACGGACATT	CAAGTCGCTT	5580
GTCTCTCTGG	CAGCTTCATC	AACTAATTTA	TCGTATCTTT	TCAAACCAAC	TTGAACTACT	5640
GAAGGGCTAT	TTGGATTATC	AAATCCTAAA	TATGTTTTTG	TAGTTTCACT	GCTAGTTGTT	5700
TTTAAAATAT	CCAGGTAAGT	AGATGGGTCT	TGATAGTCTG	GCCCCCATGA	AACTCCTCCT	5760
GATACATCCC	AATCCTCAGA	TGAAGCATTG	GCAGCATAGT	AAGTAATATT	AAGGAATTCA	5820
TCACTGTGCA	TTTGTGGAAT	ATCAACAACG	ACATTTTCAA	CACCAAGAAC	TGTTTCTACA	5880
GATTGTTTAA	AGGACTGAAT	ACGAGATATG	TAGTTTTTTG	ATGCTTGGTC	TACTGGAACG	5940
TCCAGATGAA	TAGGAAACTG	AACGCCGTCT	GCTTCTAAAG	CTTCTTAGC	TTTCGCAAAC	6000
TCTGCCTTGG	CCTTGTGAGC	ATTGAATAAA	CCATCCTGCC	CATCAGCTAA	ATTCACACCT	6060
TTCCACTCAT	CACCATAAGC	AGGAAGTTGA	GCAGCGACTA	AATCACCAA	GGTCTTCTCA	6120
CCAGCTGAAA	CAAAGTCTGG	TTTTACAAAT	AAATTACGAA	CTGCTAAAGC	TGCTCCATCT	6180
TTACCATTGA	TTTGAGCTGA	GTAAGCTGAG	CGATCAAGAG	CAAAATTCAA	GGCTTGACGG	6240
AAATCTTTGT	TAAGCAATGC	CTTCTTAGTA	GCTACTTTCT	CTGAATCTGT	AGTTTLAGAA	6300
GTATAGTTGT	AACTTTGGCG	ATCAATATTC	ACACCAGAC	CAGCAATCCC	AGAGCCTGAT	6360

755

TGTGTGTAAT	AGATATTGTC	CTTGTATTCT	TCTGCAACCT	TAGAATAGTT	GGAGCTGGTA	6420
GGGTAAAGAC	GGGCATAACT	ATAAGCTCCA	CTAGTGAAGT	TACGCTCTAG	CGACTCCTGA	6480
TCTGATCCAT	CATAGTAAGC	TAGATTGATA	GTATCTAGGT	GGACATTTTC	TTTATCCCAA	6540
TATTGCTCAT	TTTTTACAAA	CTCTACAGAA	GATTTTGCAG	TCAACCCCTT	CAACAAGAAT	6600
GGACCATTAT	AAAGCAAGGA	TGTCGGATCT	GTTGGTTTAG	CAAAATCGCT	TCCTTTTGAT	6660
GTTTCGAATT	CTTCATTCAG	AGGCCAGAAA	ATAGAATAGG	TCAACTTAGA	GTTCCAGAAC	6720
GGTTCAGGCT	GGTTCAAAGT	GTATTGTAAC	GTATAATCAT	CAACCGCCTT	GACACCAACT	6780
GTTGAAAAAT	CTGTTGAAGT	TCCTGATAGA	TAATCTGCCA	AGCCTTTAAC	CGAATTTTCA	6840
GCTAAATACA	TAGCTTCTGA	TTTTTTATCT	GCTGCGTGT	TTAAACCGTT	CACGAAATCT	6900
TTAGCCGTCA	CCTCTGCATA	TTCTTCTCCA	TCAGAGGTAA	ACCATTTAAC	CCCTTTACGA	6960
ATCTTATAAG	TGTAGGTCAA	ACCATCCTTA	GAGACTTCCC	AATCCTCTGC	AACTGCAGGA	7020
GCAAGATTAC	CGTAATTATC	GTTAGTGAAT	AAACCATCAA	TCCCATTTGA	AGTCACTACT	7080
GTTGTACTAT	TTTTACTTGA	AATCAGGTAG	TCCAAGGTTT	CTGGGTCTGC	TGTATAAACA	7140
TAGCCATAAG	CTTTAGGGGC	TGATGAATCA	GATGATTTTG	AAGAACTGCA	TGCTGCAAGT	7200
ACACCTGCTG	CTAATAAAAC	AAGACCTGCT	GTAGCAAATA	CACGATTTTT	TTTCATTTTC	7260
TACTCCTCTG	TTTATGTGAA	TTATAGATTG	ACAACCATTA	TATCACATTA	TCCATTAATA	7320
ATCAAACAAA	TTTTCAGAAAT	ATTTAGGCTT	GTTGGCACAA	ATTTTTCATT	TTTTTTGAAT	7380
ATATGATTCA	AATGTGCGTT	CGAAGTGTCA	AAGACTACAG	TGAAAATAGG	AAATTTGACG	7440
CAGAAACTTT	GGAGTTTAGG	AAGACATACA	GTAAAATGAA	ATACGGACGG	AACAATGTGA	7500
TTTTGGAATT	CAAATTAAT	TATAACAATA	TTGTAGAAGT	ATCATTCTAG	TATTCAAGAT	7560
TCAGTTTACT	ATGTCTTTTC	ACACCAACCT	TATCCCGAAT	TCAATTACTT	TTGTGATTTA	7620
CATATATAGA	TTAAGACTAT	CTTTTATACT	TTAAAATTTT	TCGCTACCTT	ATCCACTATA	7680
TGCTCCTCGC	TATCACGTTT	CTATTCATAG	CCTACGATTT	CACTATTGCT	TTCTCTGACA	7740
ATTCTTATTT	CCTGCGTCAG	ACTTAAAACG	ATCTATCCCC	AGACCATTTT	AATCCGCTAC	7800
CTCACGATAG	TCAGGCTTGG	GGAGCGCTAT	TGTATTCACC	GGTAGTGGAG	CCCTACAGAG	7860
GACTTACACC	TCAGATGCAC	GACATGCCCA	TCGTATAAAA	AATCTCCTAC	CCAAGGTAGA	7920
AGATTTCAAA	CTTATAAAAC	TTAATCCGTC	ATGTCGGATA	CCAACATTCG	ATGCTCCAAT	7980
GGAAACTGTC	ACATAACTAG	CAAGAAAATA	AAGCCTGACT	GAATCCAGAA	GAGAGCCAAG	8040
TCAAAAATTC	CGTGCACAGC	AACCACTGTA	AGGAAAGATA	GATAAAGGCC	GATAATCGGA	8100

756

CGTTTCCCCG	ACTCCTGACT	CATATCCATC	ATCAAGCGAA	CAGGAGCAAC	AGAAGACAAA	8160
ACTAATAAAA	TAGTCCCCAC	AATFCCGTAA	CTCAGAATCG	TATCAATATA	AAGACTGTGG	8220
GCATGTTTCAT	GATAAGGAGC	ATGTATCCGA	GGATAAGAGT	TCATATAGGT	CAATGGCCCT	8280
TCACCCCAAA	AAGGATTTTG	CTTAAACAAG	GCCATCCCAG	CATCCCAGAT	AGAAATGCGT	8340
TCTTCCATAG	AAGAGTCTAA	AGTACCCATF	CGAACTCCCA	AATCACTAGA	AAAGAGGAAA	8400
CTCAAACCAA	TCGCGAAGAC	CCCAATACTA	AGCCAAAAGG	CCTTCCAGTT	TTTAATAGTC	8460
GTAAGAGAT	AGATAATGTC	TCCAGCGATA	ATAGCAGGAA	AGGCAGTTCG	ATTTTGAGTA	8520
AAGTTCAAAC	CAAAGAGATT	AACAAAGCCT	GCAATCACAC	AGAATACTTT	CAACCAATTC	8580
AACTTGGTCG	TTGTAAACAG	ATAGAAAGCA	ATCATAATAC	AGAAACAACA	AATAATTCCA	8640
TAATAATTAG	GATTAAAGAA	GGTCACTTCT	GCCCGGTTCT	GATGCCACAC	CTGCATATTG	8700
GGTGAAAGAA	AAGCATAGTT	AAATTTCTTC	ACAATTTGGA	AATGTTCTAA	ACTGGCAAAA	8760
GCAGCTGACA	AGACACTACC	AAACAAGACA	AACTGCAAAA	TCAATCGAAA	GAATTTATGG	8820
GATAAAATCG	ACTGATAGTG	CAAAAAGAAA	ATAGTAAATA	GAAACATTCC	TACTGAAGCC	8880
ACAAGACCCA	TCCAATTTTG	TGCAAGAATG	GATATAACAG	TACTATAGCT	AAGAAAAAGA	8940
AGCAGCATCG	GATGCTCCCC	CATTTTCTGA	AGAATACTTT	TCATGTCTCC	TGTA AAAATC	9000
AAACTGATAA	TATATAAACA	GAGTACAACT	ACAAAAAGAT	AAAAGGGTAA	AAAGATACTC	9060
AGGATAATTC	CCAATAAAAT	CAGCTCTTTA	CTAGACAACC	CCTTCAGCTT	TTCAATAAAG	9120
CCTATTGATT	TCAAAATGAA	TCCTTTCTCT	CCAATCAGC	TGATTGAGAT	AATAGTAAGC	9180
TATCCTATAT	TGTACCACCT	TTTTAGCAAT	TTGAAAACAA	AGGAAACGTT	TTCCAAAATA	9240
AAAACCCTAT	TTTATCCACC	ATATCAAGGC	TTCAAATGA	TACTTCAACT	CCATTCTCAA	9300
TTACCCGATA	AGTCTGATTT	TGCAAATCAA	TTTCTACTAC	TGCTGTTACG	GACTTATCTT	9360
TATTTTGACG	TTTGATTACA	ATGCTGTGAG	CTGTTGGTGT	CTCTATCTCA	GTAGTCCCTT	9420
CTAGATCAAA	GGCTTCTGAA	CGGTTACGGA	AAGAAAATAG	ATGAGAAGG	GCCTTCACAA	9480
CAGGTCGTTG	CACTTCTTTT	GCTATTTCTT	CGTTGCTATA	GTAATGACGA	TTAATATTTT	9540
GACCTTCTTT	AGTTTCTTCT	AATAATTTCA	AGTCATTTCT	GCCTGCTAAT	AGACCCACAT	9600
AGTAAATCTG	AGGAATACCT	GGGGCAAAAG	CTTGAATTAG	ACGAGCGAGA	AAATACTTGA	9660
CATCATCATC	TCCAAGCGCT	GAATAGTAGG	TTGAATTGAT	TTGGTAGATA	TCTAAGTTGT	9720
TATACTCGGC	ACTAGAGTAC	TTACGTTTGA	CATTGGCTCC	AACCTTATAG	AGTTCATTTG	9780
AAGCATAGTC	AATCTCCTCA	TCGGTCAGGA	TATCCTTGAC	ATCTACTACT	CCAATCCCAT	9840
CATGGGTATC	TAGCGTCGTA	AATFGCTTCA	TCGGGCTCAT	CTTTAACCCAC	TTAGCCAAAC	9900

757

GCTCTGTTCT	GGAAGTGTAA	AGAGTATAAA	GTGTCACCAT	TGGAAGAGCA	AAATCATAAA	9960
CATAGTAATC	ATGGTCTGCT	ATTTTAAACT	GAATCGAATA	GTGTTTCATGA	ATCTCAGGTA	10020
AAAGCTCTGT	CCCATACTCA	GCAGCGATAT	CTCGAACTTT	GTCCAATAAA	TCCCAAATAT	10080
CTGGTCCAC	AAAGAAATCA	TTAGTATCCA	ATTTCTTCAC	TGCATAAGCA	AAGGCATCTA	10140
GACGAATCAA	ATCACACCCA	TTACTTGCCA	AGTGC'TGAAT	GGTCTTACGG	ATAAAT'TCCA	10200
TAGTTACTTC	TTTGGTCACA	TCAAGATCAA	TCTGCTCCTC	ACCAAAGGTA	TTCCACAAAT	10260
GTTCCACTGA	ACCATCTTCA	AACACAATCT	CTTGCTTTGG	TGCACGATCC	TTACGCTTGT	10320
AAATTAATC	TACATCAGAC	TGTGTCGGAC	GGTTTTCTGG	CCAAAAC'TTA	TCCAGTTTA	10380
AAAAGAGAGC	TTTAAAT'TCA	CTGGCTTCAT	GTTTTTCTTG	ATAGTCCTTA	TAATACTTGG	10440
ATTGACGAGA	AATATGATTA	ATCATAAAAT	CAAACATAAG	ATAATAT'TTC	TCACCTAAAC	10500
GCTTCACATC	CTCCCAATCA	CCAAAAGCTG	AGTCCACTTC	GTCGTAGTCA	ACTGGCGCAA	10560
ATCCACGATC	AACTGT'TGAT	GGGAAAAATG	GTAAAAGGTG	AACTCCTCCA	ATAGCATCTC	10620
CAAAATGCTC	TTCCAAATTA	TCATATAAGT	CTTAAAGATT	ATTTCCAAGG	CTATCAGAAT	10680
AGGTAATCAA	CATGGTTTTA	TTTTGAATTG	GCATCATTAC	TCTCCTTTTT	CTAAT'TGAAG	10740
CCAAGTCTCA	TATGATCTGG	CTTCATAAAT	AAAATTCATT	TTAAATCTCT	ATTTATCATC	10800
AAACTCGTAC	TAATATAGAC	TGTGATAAAC	AAAGTACTAC	TTTCTTGTTT	TCTGCATAGA	10860
ATTATCAACA	AGCTAAACTC	TTCTCTGTG	TCAAAGACTA	TAGAT'TCCAT	GAGCTCTTCT	10920
TATACTCTTC	GAAATCTCT	TCAAACCACG	TCAGCTTCAC	CTTGCCGTAG	GTATGGTTAC	10980
TGACTTCGTC	AGTTTCATCC	ACAACCTCAA	AACAGTGT'TT	TGAGCAACCT	GCGGCTAGCT	11040
TCCTAGTTTG	CTCTT'TGATT	TTCAT'TGAGT	ATTACTTCAC	TGCCCCGTTG	CTCATTCCTG	11100
AAATGATATG	GCGTTGGAAG	AAGAGATAGA	CAATGGTGAT	ACTGATAATG	CCGACCACGT	11160
AAGAGGCAAA	GCTTGGTCCG	TAGTCGTTGA	AATATTGGCC	TGCGTAGTTG	TATTGGAACA	11220
AAGGCAGAGT	CCACATTTTG	GAATCCCGGT	TCAAGACAAG	GAGTGGCAAC	ATGAAGTCAT	11280
TCCAGAACCA	AAGGCAT'TG	ATGATCATGG	TTGTGCGCATG	CATCGGTTTC	ATCAT'TGGGA	11340
AGATGATGCG	GAAATAGGTT	GTAAAT'TGAT	TAGCCCATC	GATCTCTGCT	GCTTCATCCA	11400
GACTTCTGCG	AATCGAGATT	TTGATATAGC	CAACATAGAG	AAAGAGGGTC	TGTGGAATCG	11460
CATAGGTCAA	GTAGAGCAAG	ATCAAACCAA	AGGTATTAGC	CAAACCGAGT	TTACTCATCA	11520
TAACCGTAAT	CGGAATCATG	ATGACTTGGG	AAGGTACGAA	GATTCGGAGG	ATTAAGAGGG	11580
TATACATGAT	GGTAAAGGCT	TTTCTTTTAC	TCATAT'TGCG	AGCGATGGAG	TAGGCTGCCA	11640

TAGGGATAAA	GATCATTACT	GCAAGTAAAG	ACAAGACAGT	GATGACGACA	GAGTTCCAAT	11700
AATAGCCTCC	AATCCCATCA	GCTAAGAGAC	GGCTAAAGTT	GTCCCATGTG	AAGTTGGTTG	11760
GAAAGCCAAA	GAAATTATCT	ACAATATCCT	TAGTGGGTTT	GAAGGAACTA	AAGAGGGTAG	11820
CAAGGAGCGG	CACTAAAATC	AGAACCGATC	CTAGAATCAA	TAGAATGTAT	TTGCCAATCA	11880
GGGCTTTTCT	TTCATCTTGT	TTCATCATGC	TTCTCCTCTT	AAATTTCAAA	TTTCTTAGAT	11940
ACTCTCAATT	GGATGATCGA	AATCACTACA	ATTAAGAAGA	ACAAGATTAC	GGCAATGGCA	12000
TTGGCATAAC	CGAATGGGT	GTTTTTAAAG	GCATAGTTAT	AAACCAAGAG	CCCAAGTGAG	12060
GTTGTGGCAT	TGTTTGGACC	ACCACCGGTC	ATGGCAAAGA	CTTGGTCAAA	GGCAGTCAGC	12120
CCACCTTTTA	GGGCTAGGAT	AAAGACCATA	GAGACTTG	GTAGCAAGTA	AGGCAATTCA	12180
ATGTTCCAGA	AAACTTGCTT	GCTAGTCGCA	CCATCAATCC	TTGCTGCCTC	TGTAATCTCA	12240
GTTGGAATAG	ATTGCAAACC	AGCTAGGAAG	ATGATGATGG	GCATAGCCAC	CCCTTGCCAA	12300
AGAAGGACAA	AGACAGCCGC	AAAGATTGCT	CCCCACTTAG	TCCCTAAAAG	ACTGGTTTGG	12360
AAAAATTCAA	TATGAAGGGC	ATTTCCAATC	GCTGGAAGAC	CGTAGTTGAA	GACTTGCTTG	12420
AAGATCAAAG	CCACTGTCAA	ACCAGATAAA	ACAGCTGGGA	AGAAGAACCA	AGCACGGAAG	12480
AAGGTTTGGC	CTTTGATTTT	AGAATTC AAG	ACACGCGCAA	TGAAGATCCC	GAGTGCAATC	12540
TCACCAACCA	CCATGGCAAT	CGCAATGATT	GCGGTAAAGC	CAATCGCATT	CATGAATTTT	12600
GGATCCATGA	AGAGGAGCTT	AAAGTTGTTT	AAGCCAACAA	ATTTGTAGTT	ATAAGTCAAT	12660
CCTGTCCAGT	TGGTAAAAC T	GTAAAAGGCT	CCTTGAAACA	TCGGCACATA	GAAGAAAATT	12720
GCTTGTAACA	AGAGGGGGAT	GACCACAAAA	GCCCATGCCC	AATATTTTTG	TAATACTTTT	12780
TTCATAGTCT	CTCTACTCCT	AATCCACATC	CGCTTTCATC	GGGTTAAAGA	AGGCATTCAA	12840
ATCATTGACC	ATGCCTTGTT	TATCACCGGT	CAAGACATAG	TTCATGGTCA	AGGTATGGAA	12900
GTCTGCTTCA	CTGGTCCAGT	ATTGTTGCAA	CCAGACCAAG	TGACGATCCG	TAAAGGCATA	12960
TTCGGTCATA	CCAGCAAGCG	GTGAATCTTC	TCCTGCTTGT	TTGACCCCTT	CGATCGCTGT	13020
TGGAGATCCG	TCCACATCGT	AGTATTTTTG	CATGACTTCT	GGACGGGTCA	TATATCCAC	13080
AAAGGCATTG	GCTTCTTTTG	GATGTTTGGT	GGTGGCTGAG	ATAGACCATG	CCAAGTCTCC	13140
CGCACCAACG	GTTAAGCTTT	GTCCTTTTTC	TTTTCTGGA	ATCATGAAGG	TCCCAATCTT	13200
AAAGTTCGGT	TTTTGTTCAT	TAATCGCTGT	GATCGCCCAA	GACCCATTTG	GTGTCATGAG	13260
GACATCCCCA	CGTGCGAAGG	CTCCGATAAC	ATCGGTATAG	CCAGCACCTT	CCCAGTTCTT	13320
TTGCTTAGAT	CCATTGATGC	GAAGGATGTC	CATGACCTTG	ATATCATCTT	TCATAATCGG	13380
ATCCGACAAT	TTAATGGCAT	TTGGTTGAGA	ATAACGAAGG	TATTGATTTG	CTTCTTTTCC	13440

TCCACCTGTT	GCTGTCGCAA	AGGCTAATTG	ATTGTAACCA	TTGAGTGTCC	AAGCATCTGC	13500
ACCTGCAATT	CAAATGGTG	TTTGTCCTTT	AGCAACGATA	TCTTTGACTA	ACTGTTCAAA	13560
TTTCATCCCAG	GTTCAGGAA	CCTTCAAGCC	CAGTTCTTCG	AATTTATCTT	TGTTGTAGTA	13620
AATTCATAA	GCATTAGCTG	TAAAAGGAAC	GTTGTAAACT	TTTTCGTTTA	CAGCATATTT	13680
TTTCAGCGTAG	CCATTTTCA	CGCGTTTCAG	GTAGTCTTTG	TTGCTCAAAT	CTCAAAAAC	13740
ACCTGCTTTT	GCCCATCTT	GCAGTTCGAT	GGACTGTGGG	TAAATATTGA	CCACATCAGG	13800
CACATCTCCT	GCGAGAACGC	GTGTCTTCAA	TACTTCACCA	GCATTTGGTA	CATGACGAC	13860
TTTGACCTTG	ATCTTAGGGT	TTCCCTTCTC	AAAATCACGA	GTGATTTCTT	CCAAGGTTTT	13920
GGTCATTTCT	TTTTTCTGGT	TGAAATACTC	GATGGTCACT	GTGCCATCCG	CAGATTTACC	13980
ATAGTTGGAG	CAAGCGCCGA	GCCCAAACAA	AGCTAAACCT	GTAGTTGCAA	GAAGTCCGAT	14040
TTTTTTATAC	CATTCATTA	GAAAGCCTCC	TTTATAAAAT	TATACACCTT	TATTTGAACTG	14100
CACCCAAAA	GTTAGACAGA	ATAAATCTAA	CTTTTGGGGT	CAGTACATAT	CATAGTTTTTC	14160
TAAAAATATA	CTGTCTACTC	AAAAAATCTC	CTTGGGATAA	GATAACAGTT	AAGCCCGCAT	14220
ACATTAGTTC	TGCACCTGAG	TAAACTTCGC	CATTTTCCTG	TAATTTATAT	AGTCCCTCTT	14280
CATCCAAATC	TTTTAATTTT	AAAGTTGTTT	CCATGGTCTC	TACAACAGAT	AAAACGCGAA	14340
CGTAGGTAC	AATCGTTTGA	TTCCCGTAAT	TAAATTGTAC	AGCTGCTTCA	TTGGATACAG	14400
TATCAGGATT	AATTAGTCTA	TACTGCTGTC	CTAACTGAAC	TACTGGTCGT	AATCTTTTAT	14460
ACAAGTTCAC	CTGATTAGCA	ATCGTAGCTT	TCTCTTCATC	TGATAAATTT	GTCAAATCAA	14520
GTTCATAGCC	CAAATTTCCC	ATCATTGCTA	CAAGGCCACG	TGTTTCTAAT	GGTGTCAATC	14580
GTCCCATCTG	ATGATTCCGT	ACTGCTGACA	CATGAGCCCC	CATAGAAATG	GTTGGATAGA	14640
GATAGGATGA	ACCGTATTGA	ATTGGTAAAC	GTGCAATGGC	ATCAGTATTA	TCACTAGCCC	14700
AGACTTGTGG	GAAATAGCGC	ATCATACCAA	GATCATTTTCG	TCCACCACCA	CCAGAGCAGG	14760
ACTCAAAGAG	AATATGGCTG	TGCTTCTCTG	TCAGATAAGA	AACGAGTTCA	TAAAGCCCCA	14820
GCATGTACTG	ATGAGATTGC	ATCTGTGTCT	CTAGATAAGT	TAATCCATTC	CCTAGCTTAG	14880
TGATATTGCG	GTTCATATCC	CATTTAATGT	AATCAATATC	ATGATAAAAT	AGGAGTTGAT	14940
CTAAGACACT	TTTCAAGTAT	TCTACTACCT	GAGGATTGGC	AAGATTAAGT	ACTAATTGAT	15000
TCCGAGAATA	AGTATGCTCA	TAGCCAGGAA	CCTGAATAGC	CCAGTCAGGA	TGTTGACGAT	15060
ACAAATCACT	ATCTACAGAA	ATCATTTTCGG	GTCTAACCA	AAGTCCAAC	TGCAAACCTC	15120
TTTCATGGAT	AGCTGAAATC	AGACTTTCTA	GACTTCCACC	CAGTTTTTCC	TCATTAACAA	15180

760

CCCAATCACC	TAAAGCACGA	TTATCATCAA	AACGATTGCC	AAACCAACCA	TCATCTAATA	15240
CAAAAAGTTC	AATGCCAACT	TTCTTAGCTT	CATCTGCTAA	CTCTAACAGT	TTTTCTCTCT	15300
GAAAGTCAAA	GTAAGTAGCT	TCCCAGTTAT	TGATTAGAAT	TGGACGTTCT	TTTTTAGAAA	15360
ATTCACCTAG	CATAATGTGC	TTCAGTACAA	AATTCTGACT	TTCATGACTA	ATACCAGTTA	15420
ATCCCTGATC	TGAATGAGTC	ACTAAAGCTA	CCGGTGTTTC	AAAGTATTCC	TCAGGAGCTA	15480
ACTTCCAAGA	AAAGTTTTCT	GGATTAATGC	CAATAGCCAC	CCGAACTTCA	TTCAATTGAT	15540
TTTTTTGAAC	AAAAGCTTCA	AAGTTGCCAC	TATACATTAG	TTGAATAGCA	AACACATTCC	15600
CAGCATCCTC	TGTGACTCCT	TGTTGCATA	GTAGAAGAGC	TGGTGTTTGA	GCATGACCAG	15660
AAGCACCTCG	GTTTGAACTA	ATCGAAAAGA	TTCCTTGTTC	TACCTGTTGA	CGTCTAACAG	15720
TCTTTTCACG	AGCATAAGCA	CCCTGCAGAG	TTACTATTTT	GTAATCTGCA	GCTGGAAAAT	15780
CAGCCATAAA	AGAAAAATCT	TTATGGATGA	CAACTTCCTG	ATTACTATTA	TTATCTAATT	15840
TACTGTAGCT	AGCAATAGTC	GCATCATTAT	TAAAAGTAGT	ATAATACAAA	GTCAGACTAA	15900
GTTGAGCCTT	AGAATCTTCT	AACATTAAGA	CAAGAGTCTC	TGTATCGTCC	ATGCTATGTG	15960
GAGAAGGTAA	GCCCTGTGGA	CCATCTGAC	CTTTTAAAAT	CTTTGCTTCT	ACAAATCGAA	16020
AGTCTGTAC	TTCAGTTACA	CTATGCTGAA	CCTGTATGGT	TGGTTTCCTA	AAATCTCCTA	16080
AGCCATGTTG	TCCAAAAATC	TGTCGCTGAG	TATCTAAACT	AAAGGTTCGA	TTAGTAGCCG	16140
TTGGATTTCC	TGAAAAGGCA	TGGTCTCGTT	CATAAACACT	ATTGGAACCT	TTATAGTTCT	16200
TAATAGTCTT	TCCTAAATGT	TTCAAAAAGTA	AGTAGCCATT	TCGATTTTCA	ATAATCAAAC	16260
TTAGATTTTT	ACTCTCAACA	TAAAATAGAT	TATTCTCTAT	CCTAACTCCC	ATTTACTTCA	16320
CCTCATCACT	TTATTGATTA	TATTTTATCA	CCTGAAATCG	CTTTCCAAA	TAGAAAAATG	16380
TCTCAAGAAT	ATGGTAAAAT	GTTAGGTAGG	AGGTAGCACA	TGTTAGTTTT	TTCAGAATAC	16440
CAGACTGGAA	CAATCGACCT	TGCCCTAAGC	TTTTATGGAT	ATGAGGAATG	CACACCTAAT	16500
TACTCTTTTG	GTCCAGCCAT	TCGTGATACA	TACGTTCTAC	ATTACATTAC	TAAAGGACAA	16560
GGAAAATTTT	ATTACAAGGG	TAAAATTGTT	GATTTAAAAG	AAGGAGATTT	CTTTCTATTA	16620
AAACCAGAGG	AACTAACCTT	TTATCAAGCA	GATAGTAAAG	AACCTTGGGC	CTACTACTGG	16680
TTAGGAATCA	CTGGAGGGAA	AGCCCTGAT	TATTTTGCTC	TTTCCCAAAT	TTCTGATCAA	16740
TCCTATCTCA	TCCAATCTGA	AACTTGTCAT	ACCCAGACTA	CTGCAAAACT	CATCTCAGAC	16800
ATTGTCCGCT	TCGCTCAGAT	TACAAAATCA	AGTGAATTAG	CTCAACTCCA	TATCATGGGA	16860
CAACTTCATG	AACTGATGTT	TCATCTGGGA	ACTATTGCTC	CCAATCAGAA	AAAAAAGAAT	16920
ATTTTCATCAA	CCCACCAACT	CTATCTTGAA	TGCAAACGAT	TAATTGATAG	CCACTATCCT	16980

761

CAATCACTTA	CAATCAAGA	TTTAGCAAAA	GAACTATCCG	TTCACAGAAG	CTACTTATCA	17040
AGCGTATTCA	AAGAATTTAA	TACCTTATCA	CCCAAAGAAT	ACCTACTCTA	CGTTCAAGTG	17100
CACCGAGCTA	GACAACTTCT	CGAAAATACC	CAAGAGTCCA	TCAAGGTAAT	TGCATACTCG	17160
GTAGGTTTTT	CAGATCCACT	CCATTTTTTCG	AAAGCTTATA	AACAATACTT	TAATCAGACT	17220
CCAAGTCATA	CAAGAAAAGA	ATACTCTCAA	TACCAACTAG	TAAGAAAGGC	AACATTATGA	17280
AATCCTACCA	AGCTGTCTAC	CAAATCCTAT	CTAAGAAAC	CGACTATATC	AGCGGAGAAA	17340
AAATCGCAGA	AAAACATATCC	CTAAGCCGAA	CAGCAATTTG	GAAAGCCATC	AAGCGACTAG	17400
AACAAGAAGG	CATTGAAATP	GATAGTATCA	AAAATAGAGG	ATATAAACTG	ATGAATGGTG	17460
ACCTTATTCT	TCCAGAGATT	CTAGAAGAAA	ATCTTCCAAT	TAAAGTCAGC	TTTAAACCCG	17520
AAACAAAATC	AACACAACATA	GATGCAAAAAG	AAGCAATTGA	TTTAGGCCAT	GAAGCAAATA	17580
CCCTCTATCT	AGCTTCCTAT	CAAACAGCAG	GCCGAGGCCG	TTTTCAACGT	TCCTTCTACT	17640
CACCACAAGG	TGGTATTTAT	ATGACACTCC	ATCTTAAACC	AAATCTCCCC	TATGACAAAT	17700
TACCATCCTA	CACACTACTT	GTAGCTGGAG	CTGTCTACAA	AGCCATTAAG	AACCTAACTT	17760
TAATAGATGT	CGACATAAAA	TGGGTCAATG	ATATCTATCT	AAACAATCAT	AAAATTGGAG	17820
GAATCCTTAC	TGAAGCAATG	ACCTCTGTAG	AAACTGGCTT	AGTCACAGAT	ATCATTATTG	17880
GAGTAGGTAT	CAATTTCACT	ATTAAAGACT	TCCCTCAGGA	ATTAAGAGAA	AAAGCTGCCA	17940
GCTTATTTAA	AGCTACAGCT	CCTATAACAA	GGAATGAATP	GATCATAGAA	ATCTGGCGTG	18000
CTTCTTTCGA	AACACCAGCA	GAAGAGCTAT	TATACCTATA	CAAAAACAG	TCATTCATTC	18060
TAGGAAAAGA	AGTCACTTTC	ACACTAGAGC	AAAAAGACTA	CAAGGGACTT	GCTAAAGACA	18120
TCTCAGAAAA	TGGAAAACCT	TTAGTTCAAT	GTGATAACGG	AAAAGAAATC	TGGCTAAATA	18180
GTGGCGAAAT	TTCTCTCAAT	AGTTGGAAGT	AAAATAACAC	AATTATAATA	TAAACGATAT	18240
AAAAATAACT	TCAGATTAGT	AATTCAATTA	AGTTTACGG	ATCTGAAGTT	TTATTGGCTC	18300
TAAAAATAAA	AAAGAGAGTP	ACAGACTCTC	ATTAAGACGG	AGAAATAAGG	ATTCTGAACCC	18360
TTGCGCCAGT	TACCCGACCT	AACGATTTAG	CAAACCGTCC	TCTTCAGCCT	CTTGAGTAAT	18420
TCTCCAATTA	ATGGGCACGA	GTGGACTCGA	ACCACCGACC	TCACGCTTAT	CAGGCGTGCG	18480
CTCTAACAC	CTGAGCTACG	CGCCCAAGTT	AAAAACTTG	GTAATTTGAA	CAAAGTTCAA	18540
AGCGGGTGAC	GAGAATCGAA	CTCGCGACAA	CAGCTTGAA	GGCTGTAGTT	TTACCACTAA	18600
ACTACACCCG	CATAAATACT	ATCAATAAAA	TGGCGCGAGA	CGGAATCGAA	CCGCCGACAC	18660
ATGGAGCTTC	AATCCATTGC	TCTACCAACT	GAGCTACCGA	GCCTTATTGC	GGGAGCAGGA	18720

762

TTTGAACCTA CGACCTTCGG GTTATGAGCC CGACGAGCTA CCGAGCTGCT CCATCCC GCG 18780
 TTAATAATAT AAAAGGAGGA TGTGGGATTC GAACCCACGC ACGCTTTTAC ACGCCTGACG 18840
 GTTTTCAAGA CCGTTCCTT CAGCCGGACT TGGGTAATCC TCCAATATTC AAATGGACCT 18900
 TGTAGGACTT GAACCTACGA CCACTCGGTT ATGAGCCGAG AGCTCTAACC AGCTGAGCTA 18960
 AAGGTCCGAC AAGATCATT TAGCGGCGAA GGGGATCGAA CCCCCGACCT CCCGGGTATG 19020
 AACCGGACGC TCTAGCCAGC TGAGCTACAC CGCCATGAAT CCGGAAGACA GGATTCGAAC 19080
 CTGCGACACC TTGGTCCCAA ACCAAGTACT CTACCAAGCT GAGCTACTTC CCGAGTTAAA 19140
 TAGAAAAATG CACCCTAGAG GAGTCGAACC TCTAACCGCC TGATTCGTAG TCAGGTACTC 19200
 TATCCAGTTG AGCTAAGGGT GCTCCATATT ATGCCGAGGA CCGGAATCGA ACCGGTACGA 19260
 TCGTTACCAA TCGCAGGATT TTAAGTCTG TCGCTCTGCC AGTTCGCCA CCCC GGCCCTC 19320
 TCTAAGCGAA CGACGGGATT CGAACCCGCG ACCCCCACCT TGGCAAGGTG GTGTTCTACC 19380
 ACTGAACTAC GTTCGCACTG TTTTCTTCTA TCTAAAAATG CCGGCTACAT GACTTGAACA 19440
 CGCGACCCTC TGATTACAAA TCAGATGCTC TACCAACTGA GCTAAGCCGG CTCATTTGTT 19500
 ATATCTTAAT GCGGGTTAAG GGACTTGAAC CCCACGCCG TTAAGCGCCA GATCCTAAAT 19560
 CTGGTGCGTC TGCCAATTC GCCAAACCCG CATATATGAC CCGTACTGGG CTCGAACCAG 19620
 TGACCCATTG ATTAAGTC AATTGCTCTA CCAACTGAGC TAACGAGTCT AAAATAACTT 19680
 GCGTTACCTT AACCGTCCG ACGGAATCGA CCCGGTAC 19718

(2) INFORMATION FOR SEQ ID NO: 100:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 4117 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 100:

CCGTGGAAAA GTCTGGATAG TGAATGGTCT TCACACAATG ACCTGAAAGA AGCCTGAGAA 60
 TAATTATGGA GAGTAGCATT CTGAGAGGTG TTAGCAGAAC CATATGACAG AGCTGTTTGA 120
 AGAGGGAATA TTGAGGAGAA AAATCCTGAG CCTACCAGTT GGAGTTGGAA AGAGCTGACT 180
 GTTAGATCAT GGTTTATTAT CCACAACCTG TGGATAACTT TGTGAATAAG AGAAGTTGCT 240
 AAAGAAGGAG ATATATAACG ATGAAGAAAA TCAAACCGCA TGGACCGTTA CCAAGTCAGA 300
 CTCAGCTAGC TTATCTGGGA GATGAACTAG CAGCTTTTAT CCACTTCGGT CCTAATACCT 360
 TTTATGACCA AGAATGGGGG ACTGGACAGG AGGATCCTGA GCGCTTTAAC CCGAGTCAGT 420

763

TGGATGCGCG	TGAGTGGGTT	CGTGTGCTCA	AGGAAACGGG	CTTCAAAAAG	TTGATTTTGG	480
TGGTCAAGCA	CCACGATGGC	TTTGTCCCTT	ATCCGACAGC	TCACACAGAT	TATTCGGTTA	540
AGGTCAGTCC	TTGGAGGAGA	GGAAAGGGCG	ACTTGCTCCT	TGAAGTATCC	CAAGCTGCCA	600
CAGAGTTTGA	TATGGATATG	GGGGTCTACC	TGTCACCGTG	GGATGCCCAT	AGTCCCCTCT	660
ATCATGTGGA	CCGAGAAGCG	GACTACAATG	CCTATTATCT	GGCTCAGTTG	AAGGAAATCT	720
TATCAAAATCC	TAACTATGGG	AATGCTGGTA	AGTTTCGCTGA	GGTTTGGATG	GATGGTGCCA	780
GAGGAGAGGG	CGCGCAAAAAG	GTTAATTATG	AATTTGAAAA	ATGGTTTGAA	ACCATTTCGTG	840
ACCTGCAGGG	CGATTGCTTG	ATTTTTTCAA	CAGAAGGCAC	CAGTATCCGC	TGGATTGGCA	900
ATGAACGAGG	GTATGCAGGT	GATCCACTGT	GGCAAAAGGT	GAATCCTGAT	AAACTAGGAA	960
CAGAAGCAGA	GCTGAACTAT	CTTCAGCACG	GGGATCCCTC	GGGCACGATT	TTTTCAATCG	1020
GAGAGGCAGA	TGTTTCCATC	CGTCCAGGCT	GTTTCTACCA	TGAGGATCAG	GATCCTAAGT	1080
CTCTCGAGGA	GTTGGTCGAA	ATCTACTTTC	ACTCAGTAGG	GCGAGGAACT	CCACTCTTGC	1140
TTAATATTC	GCCGAATCAA	GCTGGGCTCT	TTGATGCAAA	GGATATTGAA	CGACTTTATG	1200
AATTTGCGAC	CTATCGCAAT	GAGCTCTATA	AAGAAGATTT	GGCTCTGGGA	GCTGAGGTAT	1260
CTGGTCCAGC	TCTTTCGCA	GACTTTGCTT	GTCGCCATTT	GACAGACGGC	CTGAGACCA	1320
GCTCTTGGGC	AAGCGATGCA	GACTTGCCCA	TCCAGTTAGA	ACTCGACTTA	GGTTCTCTTA	1380
AAACTTTTGA	TGTAATTGAG	TTAAGAGAAG	ATTTGAAGCT	AGGGCAACGA	ATCGCTGCTT	1440
TTCATGTGCA	AGTAGAGGTG	GATGGTGTCT	GGCAGGAGTT	TGGTTCGGGT	CATACTGTTG	1500
GTTACAAACG	TCTCTTACGA	GGAGCAGTTG	TTGAGGCACA	GAAGATACGT	GTAGCATTA	1560
CAGAATCACA	GGCTTTGCCT	TTGTTGACCA	AGATTTCCCT	TATATAAACT	CCTGGATTAT	1620
CAAAAAAAGA	AGTTGTTTCAG	GAAGTAGCAT	TTGCAGAAAA	AAGCCTAGCT	GTGGCAAAGG	1680
GAGAAAATGC	CTATTTTACA	GTTAAGCGCA	GAGAATGTAG	TGGTCCTTTA	GAAGCTAAGA	1740
TTTCGATTCA	ACCGGGGACA	GGTGTCCATG	GTGTCGCCTA	TCAGGATGAG	ATTCAAGTCC	1800
TTGCGTTTCA	AACTGGTGAG	ACTGAAAAAA	GTCTGACGCT	ACCAACCTTG	TATTTGCGAG	1860
GAGATAAAAC	CTTGGATTTT	TATCTGAACC	TAACGGTGGA	TGGTCAGCTT	GTGGATCAAC	1920
TTCAAGTCCA	AGTTTCATAA	AAGAAGAACC	TTTGC CGGAT	GCAAAGGTTT	TTTTGGTTAT	1980
TAGTGACTTG	GTAACCAGCT	GAGGGTGAAA	GTTAGTTGTT	CAGCTTTTAA	GAGGTCTTGG	2040
TGTTGAATAG	TTGATACGAG	TGTTTGTCC	AGTCGGCATT	CTTTGACAAA	GTTAAAATGG	2100
TTGTGGTTTT	GTTTAGTATG	GATATCCAGC	CATTTATCTT	CTTTAGCGAG	GTAGACTCGT	2160

AGATGGTCAA AGAGAGGGAT TCCGAGGTCA TAGCTTGGTT TTCCTGGACA GGTGGATAA 2220
 AATCCGAGAG CTGACCAGAT GTACCAAGCA GAGAGACTAC CATTGTCTTC ATCTCCAGGA 2280
 TAGGCTTCCC AACTTGGGTG AAAAGCTTTC TGACGGAGCG TCTTGATAAG AAGGGCAGTG 2340
 TAGTCAGGGT AATCGCTGTA ACGGAAGAGA TAAGGAATGT GGAAACTAGG CTGGTTGGAA 2400
 ATGGCTATTT GTCCAAAAGG AGCAGTAGCC ATCTCGCTCA TTTCGTGAAT TFCGTAACCA 2460
 TAGCCTGTTG TTTCAAAGAG GGGAGCATCT TGACAGGCTT TCAAAAAGATA GTTGCTAAAAG 2520
 GTTCTTTTTC CACCCATCAG TTGATTAAG CCAGGGATGT CGTGGAGAAC GCCTAAAGTA 2580
 GCTTGAATGG CAGAGCATTG AGCGTAGTCT CGCCCCAAC TATAAGGAGA GAAGTCAGGG 2640
 TGAAAGTTTC CTTGATTGTC TCGTGCTCGC ATGTAACCTG TCTCAGCGTC AAATAGCTGG 2700
 CGGTAATTTT GTGAAGCAGC CTTGTAGGTT TCAGCGATTT CTATGTCTC TAGTTTTTTG 2760
 GCACAGCTGG CGATACAAA GTCACTATAG GCATAGTCTA GAGTATGGCT AACACTTTTCG 2820
 TGGTGGTCGG TAGAGAGGTA ACCTAGTTCT TGGTATTGGG CTAGTCCGTG GCGGCCATTG 2880
 ATGCCGAGAG GGTGCGCTTT GCTGGCTGTT TCGAGCATGG CTTGGAAGAG TTCTCCTTCT 2940
 AGGTCGGGGG TCATGTCCCT GCAGGCGCTA TCTGCGATAA TACCGTCTAA AAGTGTACCT 3000
 GGCATCATAC CCCGTTTCATC TGGAGCCAGC CATTTTGGAA GGAAACCAGT ATCGCGGTAG 3060
 CTATTGAGGA AACCTTCTAA AAAGCGTTGA TAGTGCTCCG GTATGATAAG GGCAAAGAGG 3120
 GGAAGGTGG TGCGAAGGT ATCCCAGAAA CCATTGTTGC TAAAGAGGAC ACCAGCTTG 3180
 ACAGTACCAG TAGCCAGATC CATGTGGATG GCTTGCCCTG ATTCATTAAT CTCATAAAAA 3240
 GTCTGTGGGA AGAGGAAGAG TCTGTAGAGG CAGTGGTCAA AGAAGGTTTCG GTCAGCCTCT 3300
 CCTGTCTCTA TAATGTCAA ACGATGGAGG AGATTTTCCC AATCCACTTG GGCACCTGAT 3360
 TTACAGCTAT CAAAATCTTC TTGAGGTAGA TTGATTAGAG CTTGAGAAGG AGAGATGAAA 3420
 GAAGTGGCTA GTTGCACTC GGTTFGACTA CTTGCTAAGT CAATTCGCCA GTCTCCAGCT 3480
 TCTTGGCTGA TAGCAAGAAT ATCCGTGTTT ATTTGCAGGG CAGTGAACAT CGTTAGCGAA 3540
 TTTTGTGTTAG TTTCAGTTTT ACCTTCTTGT CGCAGGGCAA GAGTCCGCTT ATCTACTTGC 3600
 TCTACTGTCA GTTCATCTGC TGCGTGAAGA TAGAGGGAGA GGGCTTTGCC TTGCTTTTGA 3660
 TTCAAACGAA TAGAAGCACC ATAGCAAGTC GGTGTGAGCT GGGTTTCAAT CTGATAACGC 3720
 AGAGAAAAGA GCTTCAAATA GTGAGGCTGG AAGCAAGCTT TATCTATATC ATAAGAAGAC 3780
 TGGCGGTGAA AGAGGCTGTC TCCCCCAGT TGAAGGTTGA CAGGTGTCAG AAGGAGCCAA 3840
 GAGTAGTCCC CAATCCAAGG ACTGGGCTGG TGAGTTAATC GAATCCCCTG AAAGATAGGC 3900
 AGATGTGGAT CAAAAACCA AGATCCATCC TGGTCACTGG TCTGGGGCAC AAAGTAATTC 3960

765

ATCCCAAAG GCACGCCTGT GTATGGCAGG GTATTTCCCC GAGAAAAGGC ATGCTTGTTG 4020
 GTAGTTCCAA AACGGGTATC GATGGTATCA AGTAGTGGTT TCATAGTCTT TCCTTTAGCT 4080
 GTTTTCTAC ATTATATCAG TAATAGAGGG CCTTTAG 4117

(2) INFORMATION FOR SEQ ID NO: 101:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 2727 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 101:

CTGGTTCAAT TATTATTCAC TCTAAGTAGT CATATGTTCT TTATTTATGT GAGTTTTTAC 60
 CTTTAAAGG ATCTTGTTAG ATGGGAGAAG GTTTAAAAG TGACAGATGA TAATACAAGA 120
 AAAGTTCGTT TATTAGTAGC CTTTTTTAGC ATTGTCATAG GCTACATCCT GAGTTCTTTC 180
 TTTATTAGCC TGTATCATTT GTGGCAAGAA GCGCTTAGAG GATTATTATG AAATCAAGAG 240
 TAAAGGAAAC GAGTATGGAT AAAATTGTGG TTCAAGGTGG CGATAATCGT CTGGTAGGAA 300
 GCGTGACGAT CGAGGGAGCA AAAAATGCAG TCTTACCCTT GTTGGCAGCG ACTATTCTAG 360
 CAAGTGAAGG AAAGACCGTC TTGCAGAATG TTCCGATTTT GTCGGATGTC TTTATTATGA 420
 ATCAGGTAGT TGGTGGTTTG AATGCCAAGG TTGACTTTGA TGAGGAAGCT CATCTTGTCA 480
 AGGTGGATGC TACTGGCGAC ATCACTGAGG AAGCCCCTTA CAAGTATGTC AGCAAGATGC 540
 GCGCCTCCAT CGTTGTATTA GGGCCAATCC TTGCCCGTGT GGGTCATGCC AAGGTATCCA 600
 TGCCAGGTGG TTGTACGATT GGTAGCCGTC CTATTGATCT TCATTTGAAA GGTCTGGAAG 660
 CTATGGGGGT TAAGATTAGT CAGACAGCTG GTTACATCGA AGCCAAGGCA GAACGCTTGC 720
 ATGGTGCTCA TATCTATATG GACTTTCCAA GTGTTGGTGC AACGCAGAAC TTGATGATGG 780
 CAGCGACTCT GGCTGATGGG GTGACAGTGA TTGAGAATGC TGC CGTGAG CCTGAGATTG 840
 TTGACTTAGC CATTCTCCTT AATGAAATGG GAGCCAAGGT CAAAGGTGCT GGTACAGAGA 900
 CTATAACCAT TACTGGTGTT GAGAACTTC ATGGTACGAC TCACAATGTA GTCCAAGACC 960
 GTATCGAAGC AGGAACCTTT ATGGTAGCTG CTGCCATGAC TGGTGGTGAT GTCTTGATTC 1020
 GAGACGCTGT CTGGGAGCAC AACCGTCCCT TGATTGCCAA GTTACTTGAA ATGGGTGTTG 1080
 AAGTAATGA AGAAGACGAA GGAATTCGTG TTCGTTCTCA ACTAGAAAAT CTAAAAGCTG 1140
 TTCATGTGAA AACCTTGCCC CACCCAGGAT TTCCAACAGA TATGCAGGCT CAATTTACAG 1200

	766	
CCTTGATGAC AGTTGCAAAA GGCGAATCAA CCATGGTGGG GACAGTTTTC GAAAATCGTT		1260
TCCAACACCT AGAAGAGATG CGCCGCATGG GCTTGCATTC TGAGATTATC CGTGATACAG		1320
CTCGTATTGT TGGTGGACAG CCTTTGCAGG GAGCAGAAGT TCTTTCAACT GACCTTCGTG		1380
CCAGTGCGGC CTTGATTTTG ACAGGTTTGG TAGCACAGGG AGAAACTGTG GTCGGTAAAT		1440
TGGTTCACCT GGATAGAGGT TACTACGGTT TCCATGAGAA GTTGGCGCAG CTAGGTGCTA		1500
AGATTCAGCG GATTGAGGCA AGTGATGAAG ATGAATAAGA AATCAAGCTA CGTAGTCAAG		1560
CGTTTACTTT TAGTCATCAT AGTACTGATT TTAGGTACTC TGGCTCTAGG AATCGGTTTA		1620
ATGGTAGGTT ATGGAATCTT GGGCAAGGGT CAAGATCCAT GGGCTATCCT GTCTCCAGCA		1680
AAATGGCAGG AATTGATFCA TAAATTTACA GGAAATTAGG CTGGAGAACC AGCCTTTTTC		1740
TAAAGATAAG GAGAAATATG AACAAAAAAA CAAGACAGAC ACTAATCGGA CTGCTAGTGT		1800
TATTGCTTTT GTCTACAGGG AGCTATTATA TCAAGCAGAT GCCGTCGGCA CCTAATAGTC		1860
CCAAAACCAA TCTTAGTCAG AAAAAACAAG CGTCTGAAGC TCCTAGTCAA GCATTGGCAG		1920
AGAGTGCTTT AACAGACGCA GTCAAGAGTC AAATAAAGGG GAGTCTGGAG TGAATGGCT		1980
CAGGTGCTTT TATCGTCAAT GGTAATAAAA CAAATCTAGA TGCCAAGGTT TCAAGTAAGC		2040
CCTACGCTGA CAATAAAACA AAGACAGTGG GCAAGGAAAC TGTTCCAACC GTAGCTAATG		2100
CCCTCTTGTC TAAGGCCACT CGTCAGTACA AGAATCGTAA AGAACTGGG AATGGTTCAA		2160
CTTCTTGGAC TCCTCCAGGT TGGCATCAGG TCAAGAATCT AAAGGGCTCT TATACCCATG		2220
CAGTCGATAG AGGTCATTTG TTAGGCTATG CCTTAATCGG TGGTTTGGAT GGTTTTGATG		2280
CCTCAACAAG CAATCCTAAA AACATTGCTG TTCAGACAGC CTGGGCAAAT CAGGCACAAG		2340
CCGAGTATTC GACTGGTCAA AACTACTATG AAAGCAAGGT GCGTAAAGCC TTGGACCAAA		2400
ACAAGCGTGT CCGTTACCGT GTAACCCTTT ACTACGCTTC AAACGAGGAT TTAGTTCCTT		2460
CAGCTTCACA GATTGAAGCC AAGTCTTCGG ATGGAGAATT GGAATTC AAT GTTCTAGTTC		2520
CCAATGTTCA AAAGGGACTT CAACTGGATT ACCGAACTGG AGAAGTAACT GTAACCTAGT		2580
AAAAGATACG CCTACACTCC TATGTCACCT ATGGATGTAG GAGTTCCTTT TACTAGTTTA		2640
AGCAGGACTA AGACAGGTAC TAAGACAAAA TAGCAACTTC TAAAATAAC TTCCAGTTTT		2700
GGGAGAGAGA TGAAGTTAC TTTGAGA		2727

(2) INFORMATION FOR SEQ ID NO: 102:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 5717 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 102:

TTTTTTGTAG ATTTAAGTGG GGTGCAATTC CTAAAAAATA AAAAACAATT TTGAAAATT	60
ATGTTAGCAG GAATTGCTTC AAATTCGATT TTATCACTTA CAGGTTTACT TGTTTTATTG	120
TTCACATCGT ATAAATTGCT TGGACTCTTA TTTTTTATCA TTAACCTAGG TATGATTTTT	180
ATTAATCAA TTCCTTTTT TCAGTATGAT AGTGGTATTA TTTTAAGATA CTTGAATTCT	240
AACAATAATA ACTTGAATTT TCAATATATA GTTCAACTTT TAATAGCATT TGTTATTATT	300
TATTTTCCTT TGAGTCAACT ATTACAGTTT TTGACACCCA ATATTATTGT TCGTAGTATA	360
GGAGGGTGG TTGTTTCTAT ACTGCTTCT ATATTATATA TGATAGGAAG GACGAAATAT	420
GTTCTACGTA AATAGTTATG TTTTGTCTTA TAAAAAGAA GGTATAATGT ATTTACGTGG	480
TCGGAGTATG CGGAAATAG CTATAGAACC TCAAATTCG CAAGAATTTA TCAACGATCT	540
ATTTAATAGT TGTAAGGAAC TATTAGAGAT AGAAGAAGTA TTAGGCAGTA AACTAACATT	600
TGAACTATAA ATGAACAAAT TTTAATTCG GATGAGATAG ATATTGATAG TAGATATFCT	660
AGAACTAAAG GTTACTATTC GTTATTTTAT AATGAAGAGT ATAATAAAAT ACAGAATAAA	720
ACAGTATTAG TATTAGGAGC AGGAGTCTTA GGATGTTATA TATCTCTAAG TCTAAGTATG	780
TATGGAGTGA GGAAACTTAT TGTCGCTGAT TACGATATAA TAGAACCATC AAATTTAAAT	840
AGGCAAATTC TTTATACAGA GTCGGATGTT GGTAAGGAGA AGATTAATGT TCTTTCTGAA	900
AAAATACACA AGTATAATTC AGATGTTTCAG GTAGTACCTA TTTCTATTAA AGTTTCTTCA	960
GTAGAAGAAT TAGAAAAAAT TGTTGCGGAA TATGGGAGTA TAGATTTTAT CGTTAAAGCA	1020
ATTGATACGC CCATTGATAT TATAAAAATT GTCAATCAAT TTGCTGTATC GCATAAGATA	1080
TCCTACATAT CAGGAGGGTT TAATGGATGC TATCTTATTA TTGATAATAT ATATATCCCT	1140
ACCATCGGTT CTTGCTTTGG TTGTCGGAAT ATAAACAAAG ATATAAATAA GTACACTTTA	1200
TCTGATAAGA CAAAGTGGCC GACTACACCA GAGATGCCTG CTATTTTGGG AGGGATAATG	1260
ACTAATTTAA TAATTTAAAT ATTTCTGGGA TGTTATAATG AAATCCTAAT AGATAACGCT	1320
TACGTTTATA ATATGAGAAA TCATGCTCTA AGTCAAGAAA AATATGTCTT GGAAAACGGA	1380
GAATGTCCAA TTTGTAAAAA AATAATAAAG TGAAAGATAA CAATATTAGA GCGAAAACAT	1440
TTATTCGTTT AGTTTGTTTT TGCTTATTAT CAGGAGGAGT AGCTTTTTTA TCTGCTATTG	1500
GGCAGTTCAC TGTTATAGAA ACACAATTAA TAGTATTGTT CTTGGGTATT ATTTTTGCTA	1560
TATATTATGC TTACTIONAAT AAAAATATTC AAACATCATT GGAAAATATA GTATGGCTTT	1620

TTTCATCGTT	TGAGATTTTA	TTTTTGCTTG	TAAATTTTAG	AACATTTATT	CAGTTACCAG	1680
TGGATATTTT	TATTGGTATG	ATAATATTTT	TAATGCTGTG	GATATTTATT	ATGTTAGGTA	1740
TAGTGTGTCT	TAGTTATTAT	ATAACTTTAT	TATTTAGCAA	GGAGGCTTAG	TATGTTTAAA	1800
AAAATAGGTA	TAATGAGCAT	TTGCATATAT	ATAATTATTT	TATACTGCTT	GAGAATGTAT	1860
CGTATTATCA	ATAATATTGA	AACAATCTTG	CTAACGGTTA	TATGCTTAAT	GTTATTGTTT	1920
TTTTTAAGAC	GTTTATTTGA	TAAAGATAAG	TAAATAGATG	TTAAGTAAAA	ATGTAGAATA	1980
TAAAGGAGGT	GCAATGAGTA	TGATTGAAGT	TAGCCATTTA	TCAAAAAGTT	TTGGTGATAA	2040
AATAGCTTTA	AATAATATAA	GCTTCACTGT	TAAAGAAGGT	TAGATTTTTG	GATTTTTTAGA	2100
ACCATCTGGT	TCTGGAAAGA	CCACAACGAT	TAATATTCTG	ACTGGGCAGT	TCCTTGCCGA	2160
TAAAGGACAA	TCTATTATTT	TGGGACAAAA	ATCTCAAAAAT	TTAACAAGCG	GTGAATTTAA	2220
GAGAATTGGA	TTGGTTAGCG	ATACAAGTGG	ATTTTATGAG	AAAATGTCTC	TGTATAACAA	2280
TCTTCTTTTT	TATAGTAAAT	TTTATAATAT	TAGTAAATCA	CGTGTTGATA	ATTTGTTAAA	2340
GCGAGTAGGA	TTATATGATA	GTCGCAAGAT	GGTAGCAGGA	AAATTATCCA	CTGGAATGAG	2400
GCAACGAATG	CTTTTAGCAC	GAGCTCTTAT	CAACAACCCC	GCTGFACTCT	TTCTGGATGA	2460
ACCGACCTCA	GGTCTAGATC	CCACAAC TTC	TCGAACAATT	CATGAGTTAA	TTTTAGAATT	2520
GAAAACAGCA	GGGACAACGA	TTTTTCTAAC	GACTCATGAT	ATGAATGAAG	CAACTCTTTT	2580
ATGTGATTAT	GTTGCCTTAT	TAAATAAAGG	GAAATTAGTT	GAGCAAGGAG	CTCCTTCTGA	2640
ACTCATTCOA	AGATATAATA	AAGATAAAAA	GATTAAGGTT	ACAGATTATA	ATGGGAATCA	2700
GATAACTTTT	GATTTTACAT	CACTAGAACA	GGTATCTCAG	ACTGATCTGG	AAAATATTTT	2760
TTCAATTCAT	TCATGTGAGC	CTACTTTAGA	AGATATTTTT	ATCACATTAA	CAGGAGGAAA	2820
GCTAAATGCT	TAAACGGTTT	CTGGCTTTGG	TATGGTTGCG	TTGTCAAATC	ATCCTTTCCA	2880
ATAAGAGTAT	TTTATTGCAA	GTTTLAGTGC	CTTTTGCTTT	CACATATTTT	TATAAATATC	2940
TTATGGAAAC	ACAGGGGAAG	GTCAACGATC	AACAGGCATT	AGTTCTTTTG	ATGATGTGTF	3000
TACCTTTTTC	TTTTTCTTTG	GCTGTGGGAA	GTCCTATAAC	TATTATCTTG	TCTGAAGAAA	3060
AAGAAAAGTA	CAATTTACAA	ACTCTTCTGT	TGAGTGGTGT	TAAAGGCTCC	GAATACATTT	3120
TATCAACTAT	GTTTCTTCCT	TTTTTGCTAA	CTTTTGTGAT	TATGGGAACT	ACTCCTCTTA	3180
TTTTTAGGAGT	TACAATTGTA	CATACTTTTA	ATTATATTAC	AATCGTTCCT	CTAACCTCTT	3240
TATCCATCAT	TTTATTCTAT	TTATTGATAG	GTTTAAACCG	GAAGAGCCAA	GTAGTAGCTC	3300
AGGTATACAG	TCTTCTGCT	ATGATTTTAG	TTGCTTTCTT	ACCGATGCTA	TCTGGTTTGG	3360
ATAAGACAGT	TGCGAAGATA	ACAGATTATA	GTTTTATGGG	ACTATTTACT	AAGTTTTTCA	3420

CAAAATGGGA	GGAATTTTCA	TGGAATAAAA	CTCTAATTCC	TAATCTAACA	CTACTTATTT	3480
GGATTGTTCT	TCTATTAACT	TTAATTACGA	TAACTATTAG	GAAAAAGAAA	ATTTCTTAAT	3540
TGAGTTATTT	TAATGATTAT	AAACACAAGT	GGGAAGGAAA	AAATGAACTG	ATCTTTTGA	3600
CAGCAATTC	ACAGAATAGT	CTTATTGCTA	TATTTTGATT	TGAGTGTACG	AAAAAAGAAA	3660
AATAACAATA	GTGCTCATA	TAATTGCAGA	AGTTTGGGT	GATAAGATAA	CTGATAAATT	3720
GCAATAAAAA	ATGCAACATT	TTTAAATCTC	CTCTATAAGT	GCTTCAAAAA	GTGCTTCAAA	3780
ACCTGTCTTG	TAATCCAAGT	ATTTTGGGG	ACGGTGATTA	ATAAGCTAGC	AAAGCATCAT	3840
TAAGATTTT	TTCGGTAAT	GTTGCCAAAT	CGGTTAAGA	AAATACTCAC	GAAGAAGTCC	3900
ATTGCGATTC	TCATTACTTC	CCCTTGCCA	AGATGAATAG	GCATCCGCAA	AATAAACACAG	3960
AATCCCAT	TGTTCAATTA	AAGGTAACA	AGCAAACCTC	TTTTCTCTGT	CCGAAGTGAA	4020
AGTCTTTAAC	TATTCTTTTG	GAAAGAGTCT	TGTGAGGTGT	TCAATAGCAG	TCAACATGGA	4080
TTTAGCTGTT	TTTACTTGAC	AAGTGCTAGT	AGAAATAATA	GAATAGTAAA	AAACCTTTAA	4140
AGCAGTCCAG	AGAGGCAGCT	AAGGTTAGAC	GGTAAAGGG	TGGAGACTAC	CCATTTTTCG	4200
TGGAACCTTG	CTGTTGGCAG	GTCCTTTTT	TCGTGGCTTC	TGTTGGCCAG	ACTCTCTCAC	4260
TAGTAAAGGT	AAAAGGAGAA	ACCTATGCGA	GAACATCGTC	CAATCATTGC	TCTTGATTTT	4320
CCTAGTTTTG	AGGCGGTCAA	GGAATTTTTA	GCTCTTTTCC	CAGCAGAAGA	AAGCCTTTAT	4380
CTCAAGGTAG	GGATGGAGCT	TTATTACGCA	GCGGGCCCTG	AGATTGTGTC	CTACTTAAAA	4440
GGTTGGGTC	ATAGTGTCTT	TTTGATCTC	AAACTTCATG	ACATTCCTAA	TACAGTCAAG	4500
TCAGCCATGA	AGATCTTGTC	TCAGCTTGGT	GTCGATATGA	CTAATGTCCA	TGCGGCTGGT	4560
GGTGTAGAGA	TGATGAAGGC	GGCGCGTGAA	GGTCTGGGA	GTCAAGCCAA	ATTGATCGCT	4620
GTAACCTCAGC	TCACATCAAC	GTCAGAAGCT	CAGATGCAGG	AGTTTCAAAA	TATCCAAACC	4680
AGTCTGCAAG	AGTCTGTGAT	TCACTATGCC	AAGAAGACAG	CTGAAGCTGG	CTTGGATGCT	4740
GTTGTTTGCT	CGGCTCAGGA	AGTACAAGTC	ATCAAGCAGG	CTACCAATCC	AGATTTTATC	4800
TGTCTGACAC	CAGGGATTCG	TCCAGCTGGT	GTTGCAGTTG	GAGATCAAAA	ACGAGTCATG	4860
ACACCTGCTG	ATGCCTATCA	AATCGGCAGT	GACTATATCG	TAGTGGGACG	TCCCATTACC	4920
CAAGCTGAGG	ATCCTGTTGC	AGCTTATCAT	GCCATCAAGG	ATGAATGGAC	ACAGGACTGG	4980
AATTAAGAA	CTAGATTAGA	AAAATAAAG	GAGAATACCA	TGACACTTGC	TAAAGATATC	5040
GCTAGCCACC	TCTTGAAAAT	CCAAGCCGTT	TACCTCAAAC	CAGAGGAACC	CTTCACTTGG	5100
GCATCTGGTA	TCAAGTCACC	GATTTACACT	GATAATCGTG	TGACACTAGC	CTATCCAGAA	5160

770

ACTCGTACCC TAATTGAAAA TGTTTTTG TG	GAAGCTATCA AAGAAGCCTT TCCTGAAGTA	5220
GAAGTGATTG CAGGAACTGC AACAGCAGGG	ATTCCACACG GAGCCATTAT TGCTGATAAG	5280
ATGGACTTGC CTTTTGCCTA CATCCGTAGT	AAACCAAAAG ACCACGGAGC TGGTAATCAA	5340
ATCGAAGGTC GCGTAGCTCA AGGTCAAAAA	ATGGTAGTGG TTGAAGACCT TATTTCAACG	5400
GGTGGTTCAG TTCTTGAAGC TGTAGCAGCA	GCCAAGCGAG AAGGAGCAGA TGTAAGTGG	5460
GTTGTAGCGA TTTTCAGCTA CCAATTGCCA	AAAGCAGATA AGAACTTTCG AGATGCTGGT	5520
GTAAACTTG TGACGCTTTC AAATATAGC	GAGCTTATCC ATCTAGCCCA AGAAGAAGGT	5580
TACATCACGC CAGAGGGCCT TGATCTTCTA	AAACGCTTTA AAGAAGACCA AGAAAATTGG	5640
CAAGAAGGTT AGGTCAGTAA GATAAAGAGA	GACGAGGCTA CCGAGTCTCT TTTACCATTT	5700
TATTTAAAAAT ATGACAG		5717

(2) INFORMATION FOR SEQ ID NO: 103:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 5558 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 103:

CCTGGACTTT CTAAAATGAA ATCTTGCGAC	CTGGATCAAG CCCTTCATGA GCATTTTTCA	60
GAAGAAGAAT TAGCTGGTCA CTTTCATGTC	CTTCTATGGA CTTTTTTTAC AATGGCATTG	120
CTATCACACC CAATACCTAT CTAAGCGCCT	GGTTCGTAAA CTTTATTGCA GCTCTTCCTC	180
TAAATTTCTT AATTGTTGAA CCAATTGCCC	GTTTTTATACT AAGTTCTTTT CAGAAACCAT	240
TTACTGGGGA AGAAGTTGAA GATTTTCAAG	ATGATGATGA AATCCCAACT ATTATCTAAG	300
CCAGTTCTGT AAACACTATA TATTTGAAAT	CCACTTCTTT TTAGGGTGCA ATGGTTATAA	360
ATGAATTTTT GAGAGGATCA GAATGAAAAA	ACTAGCAACC CTTCTTTTAC TGCTACTGT	420
AGCCCTAGCT GGGGTAGTCA GCGTCCAACG	CAGTCTGCGT GGTGATGATT ATGTTGATTC	480
CAGTCTTGCT GCTGAAGAAA GTTCCAAAGT	AGCTGCCCAA TCTGCCAAGG AGTTAAACGA	540
TGCTTTAACA AACGAAAACG CCAATTTCCC	ACAACATATCT AAGGAAGTTG CTGAAGATGA	600
AGCCGAAGTG ATTTTCCACA CAAGCCAAGG	TGATATTCGC ATTAAACTCT TCCCTAAACT	660
CGCTCCTCTA GCGGTTGAAA ATTTCTTCAC	TCACGCCAAA GAAGGCTACT ATAACGGTAT	720
TACCTTCCAC CGTGTTCATCG ATGGCTTTAT	GGTCCAAACT GGAGATCCAA AAGGGGACGG	780
TACAGGTGGT CAGTCCATCT GGCATGACAA	GGATAAGACT AAAGACAAAG GAACTGGTTT	840

CAAGAACGAG ATTACTCCTT ATTTGTATAA CATCCGTGGT GCTCTTGCTA TGGCTAATAC	900
TGGTCAACCA AACACCAATG GCAGCCAGTT CTTTCATCAAC CAAAACCTCTA CAGATACCTC	960
TTCTAAACTC CCTACAAGCA AGTATCCACA GAAAATTTATT GAAGCCTACA AAGAAGGTGG	1020
AAACCCTAGT CTAGATGGCA AACACCCAGT CTTTGGTCAA GTGATTGACG GTATGGATGT	1080
TGTGGATAAG ATTGCTAAGG CCGAAAAAGA TGAAAAAGAC AAGCCAACCTA CTGCTATCAC	1140
AATCGACAGC ATCGAAGTGG TGAAAGACTA CGATTTTAAA TCTTAAAAAC CAAAAAATA	1200
CAGTATCCAC ATTCGGTACT GTATTTCTTT TACTCTCATT CTTAAGTTAA ATTATTAAAA	1260
TCCCATATTT GGTCTATCCA GCCTTCATAA AAGTCTGGCT CGTGGCAGAC CATAAGGATA	1320
GATCCCCTAT ATTCTTTGAG AGCGCGTTTG AGCTCATCCT TTGCATCCAC ATCCAAATGG	1380
TTGGTCGGCT CGTCCAGCAC TAAAACGTTG TTTTCACGAT TCATCAAGAG ACAGAAACGA	1440
ACCTTGGCTT GCTCTCCCC TGATAAFACT TGAATCTGGC TTTCAATATG TTTGGTTGTC	1500
AAACCACAAC GGGCAAGGGC TGCACGGACT TCTGCTTGAT TAAGGGCAGG AAAGGCATTC	1560
CAGACAGCTT CAAGAGGAGT TTGGCGATTA CCGCCTTCTA CTTCTGCTC AAAATAACCA	1620
AGTTCTAAAT AATCTCCACG CTCCACTTCC CCAGCGATTG GCGAGATAAT GCCCAAGAGA	1680
CTCTTCAAGA GAGTTGTTTT TCCAATACCA TTAGCACCAA TAATCGCAAC CTTTTGATTG	1740
CGTTCGAAGG TAAGATTTAA AGGCTTAGTA AGAGGACGGT CGTAACCAAT TTGCAAGTTC	1800
TTGGCTTGGA AGATAAAGCG CCCTGGTGTA CGAGCTGGTT TGAAATCAAA GGATGGTTTT	1860
GGTTTCTCAC TTTGGAGTTC GATAATATCC ATCTTATCCA ATTTCTTTTG ACGAGACATA	1920
GCCATATTAC GAGTTGCAAC ACGGGCTTTA TTACGAGCCA CAAAGTCCTT GAGGTCTGCA	1980
ATCTCTTTCT GCTGGCGTTC GTAGGCTGCC TCTAGCTGAG ATTTCTTCAT AGCATAAACT	2040
TCTTGGAACT GGTAGTAGTC ACCAGAGTAA CGCGTCAGCT GTTGATTTTC CACATGATAG	2100
ACAATATTAA TAACGTCATT GAGGAATGGA ATATCGTGCG AAATGAGAAC AAAGGCATTC	2160
TCATAGTTTT GGAGATAGCG CTTGAGCCAA TCAATATGCT CAGCATCCAA GTAGTTGGTC	2220
GGCTCGTCCA ACAGCAAGAT ATCAGGCTTT TCAAGGAGAA GTTTTGCCAA AAGCACCTTG	2280
GTTCTTTGCC CACCTGACAA AGAAGTTACA TCCGTATCCA TGCCAAAGTC CATAACACCA	2340
AGAGCACGCG CTACTTCGTC AATCTTAGCA TCCAAGGTAT AGAAATCACG ACTCTCCAGA	2400
CGGTCTTGAA GTTCTCCTAC TTCTTCCATG AGAGCATCAA CATCCGCGCC GTCTTCAGCC	2460
ATTTTCATAT AGAGGTCATT GATACGAGCT TCAGCTTTGA AAAGCTCATC AAAAGCCGTA	2520
CGGAGAACAT CACGCACCGA CTGTCTTTCA GCAAGGACAG AGTGCTGATC CAAGTAACCA	2580

772

GCCGTCACAT	ATTTGGACCA	CTCAACCTTT	CCTTCATCTG	GCAGCATTTT	ACCAGTCACG	2640
ATACTCATAA	AGGTTGATTT	TCCTTCACCA	TTGGCACCGA	CCAGGCCGAT	ATGTTCTCCC	2700
TTGAGGAGAC	GGAAGGACAC	ATCTTCAAAA	ATTGCACGGT	CACCAAAACC	GTGACTCAGA	2760
TTTTTAACTT	CTAAAATACT	CATTTTAATT	CCTTACCTTG	TTTTTATGTA	ATCGTTTATA	2820
AAGGAGCCAA	GCCAGATAGC	CACCCAAAGT	GTTGGTCCAC	AAATCATCAA	TCTCAAAGAC	2880
GCGATTGAAA	TCAAAGAAAA	AGTCCAAGAT	TAATTGCGTA	CACTCGATTC	CAAGACTCAC	2940
AAGAAAACTA	AAAAGAAGGA	CCTTTTGTGT	TTTCCGCAA	TTTGAAATA	GATAAAGGAG	3000
TTGAAAAATC	AGAGGAAAAA	ACAAGAAGAC	ATTGAGGATA	TTTTGTAAAA	AAATCCAACA	3060
TAATTGTCCA	ATGTCACTCA	CTTCGCCAG	TTTCCAGAGA	GAATTGAAAG	GAGTCAAAAG	3120
AAAAACCAGG	CGTCCAAGAT	GCTGAATACC	TGGAGTCCC	ACTCCCACGG	TAGATTGTTC	3180
TTGAGGAGTA	AAGCAAAAAC	AGACAATGCA	AATGCTATAG	AAAATGACTC	CCCAGACCAA	3240
AATATGATTA	TAAGTCTTCT	TCATCATTAA	GGATTTACCG	CTGCGACTGC	CTTCTGGCGG	3300
TCACGTTTCA	TTGTGTTAGA	GCGCAATTGT	CCACAAGCTG	CGTCAATATC	TGTACCATGC	3360
TCTTGACGAA	CCACACAGTT	GACCCCTTTT	TTCTTAAGCG	TATCATAGAA	AGCCAACACG	3420
CACTCTTTGG	GACTACGGCT	ATATTGGTCA	TGCTCACTAA	CTGGGTATA	AGGAATCAAG	3480
TTTACATAAG	ACAATTTCTT	GATGTTCTTG	AGCAATTCAG	TCAATCCAA	GGCTTGTCT	3540
ACACCGTCGT	TGACTTCATT	AAGCATGATA	TATTCAAAGG	TTACACGACG	GTTTGTGTTC	3600
TCAATGTAGT	ATTCAAATAGC	AGCAAAGAGT	TTTTCAATCG	GAAAGGCACG	GTTAATCTTC	3660
ATGATACTTG	AACGAAGTTC	ATTGTTAGGT	GCGTGAAGAG	ACACGGCAAG	ATTGACCTGA	3720
ACCCCTTCAT	CAGCAAAGTC	ACGAATTTTA	TGAGCCAAAC	CTGAGGTGA	AACCGTGATG	3780
TGACGAGCAC	CGATAGCCAT	TCCTTTATCA	TCATTGATAG	TACGAAAGAA	ATTCAAGACA	3840
TTGTTGTAAT	TATCAAAGGG	CTCACCGATT	CCCATGACAA	CGATATGGCT	GATGCGTTCA	3900
TCCTGACCAC	GTCATCAAAA	GTATTCTGA	ACCAGCATGA	TTTGCCTAC	GATTTCACCG	3960
TTATTGAGGT	CACGTTGCTT	CTTAATCAAA	CCAGAGGCAC	AGAAGGTACA	ACCGATATTA	4020
CAGCCGACCT	GAGTGGTCAC	ACAGACAGAT	AAACCATAGT	GTTGACGCAT	GAGTACAGTC	4080
TCAATTAACA	TACCGTCGGG	CAATTCAAAG	AGATATTTGA	CTGTACCATC	AGCAGACTCT	4140
TGCACAATAC	GTTGTTCAA	GGGATTGACC	ACAACTGGT	CATTGAGCTT	AGCAATCAAA	4200
TCCTTGAAA	GGTTGGTCAT	TTCTTCAAAT	GACTGCACAC	GTTTACGGTA	GAGCCATTC	4260
CAGATTTGAT	CTGCACGGAA	TTTCTTTTCT	CCCTGCTCCA	ATACCCATTC	CTGCATGGTT	4320
TGATGTACCA	AACTATGAAT	TGAGGGTTTC	ATTTCTTCTC	CTTATTCTCT	ACTCACTTCT	4380

773

GACGAATGAC AAAATGACGT TGTCCCTTGT CGTCTTCTG ACGACGTCTA TTTTCTTAT 4440
 CTGCATTCGA CTTTCGTTTA GTTGTGAGTCG GTTTCCTTCC TTTTCTAGAA GGTGTTTCTT 4500
 CTTC CGTCTT ACGCATTTTC TTGTCAAATG ATGCTCGCTT AGGGGCTTCA TTTTCTAAGA 4560
 CAAAATAGGC ACAACCATAA CTACAATACT CTAAAAGGTA GTCTTGTAAG CACTGATTT 4620
 TTTC AAGTTT TTCTTCTGTT CGGTCATCCT TGTAAAAACC TCGTAGGCGA AGCTGTTCTG 4680
 TGCTCCAGTC CCCCACGATA TAATCAAAC TGGTTAATAC TTCTGAAAA CGCTGATTAA 4740
 AAGTCGTAC ATCAAAGGCA TCCTTGATAT TTTCAACCA GAAAAAGCT ATCCCTTCCG 4800
 TTTGACCTT GTCCCGTGT AAATGGAAC CCGGACCAG AACTTGTTA TAGTTGTATA 4860
 ATTCAGGTGC AATTTCTTTT CGCATAGATA TCCTTTTTTC ACGATTACTT AATACTTTAT 4920
 TCTACCATAA TTTCTAGCAG TTAGCACGTT TCTCATAAAA ATGAAAAAAG TCTGACGATT 4980
 TTGTCAGACC AGAATCTTAT AACCTAAAA GAGAAGAACA ATCTTCCCT CCAACTATCA 5040
 TTATTTAGCA GCTGCGTACA ATTCATCTAC TTTATCCAG TTGATTACTG AAAAGAAAGC 5100
 TTTGATGTAG TCAGGACGCA CGTTGCGGTA TTTACAGTAG TAAGCATGTT CCCAAACGTC 5160
 CAAGCCCAAG ATTGTTTTT TACCTTCTGA GATTGGTGTG TCTTGGTTTG CTGTTGAAGT 5220
 CACTTCAAGT TTCCTTCTT TGTGACAAC CAACCATGCC CAACCTGAAC CAAAACGAGT 5280
 TGTGCTGCT GCAGTGAAG CTGCTTGAA TTCTTCAAAT GAACCAAATG TTGCATCGAT 5340
 TGCTGCTGCC AGTTCTGCTG AAGGAGCTGT TTTCTCGGGA GTCATCAATT CCCAGAAAAG 5400
 AGCGTGGTTC AAGTGTCGC CACCATTGTT GATAAGTGCT TGACGGATAT CAGCTGGGAT 5460
 AGATTCTACA TCAGCAAGCA AGGCTTCAAG GTCTTCACCG ATTTCAGGGT GTTTTCTAA 5520
 AGCTGCATTG GCATTGTTGA CATAAGTTG ATGGTGT 5558

(2) INFORMATION FOR SEQ ID NO: 104:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 6735 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 104:

GGAATTGTAA ATATCATATT GTTTTGCAC CCAAATATCG TCGTCAAATC ATTTATGGCA 60
 GATACAAAGC TAGTATCGGA AGAATCATA GTGACTTATG TGAGCGTAAG GGTGTAATAA 120
 TCCATGAAGC GAATGCTTGT TCAGACCATA TTCACATGCT TATCAGTATT CCTCCGAAAC 180

774

TTAGTGTTC	GTCCTTATG	GGCTATTTAA	AGGGCAAGAG	CAGTTTGATG	ATTTTGTATA	240
AGCATGCGAA	TTTAAAATAC	AAATATGGCA	ATCGCAAGTT	TTGGTGTAGA	GGCTATTTATG	300
TAGATACGGT	AGGCCGTAAT	CAGAAAAGTA	TAGCTGAATA	TATTCAGAAT	CAATTACAAG	360
AAGACAGAGT	AGCAGACCAG	CTCACGTTAT	TCGAGTCAGT	AGATCCGTTT	ACTGGCGAAA	420
TAAATAAGAG	GAAGTAACTA	AGGTGCTTTA	GCACCTGCTC	GGGAAAGTGG	TGCGCGAGGA	480
AGCTATTTTCG	GTGGGCCTTT	GGCCCTGGCC	GGTAGAAGCG	GCTTATAGCC	GCAGAACAAA	540
CCACCAGTTC	ACACTGGTGG	TTTTGATTTA	AAAACTTGA	TACATAAAAA	TAAAAGTCTA	600
TATAAAGGAT	GGTAAAATTC	CTGTTGTCCG	ATTTGGACAA	TATCCTAAAT	AGTTACAATA	660
TATGGTCTAT	ACTTTTCTTT	AGGAGAAAGC	TAGATGTACA	GACGTTTGAG	AGATTTGAGG	720
GAGGATCATG	ATCTGCCCCA	AAAGCAAATA	GCTACAATAC	TTTCGTTTAC	AAATTCAGCT	780
TATGCCAAAA	TTGAACGGGG	TGAGCATGCG	TTGACGGCTG	ATGTATTGGT	TAAACTCTCA	840
GATTTCTATG	ACGTCAGTAC	AGACTATTTA	TTGGGATTAA	CTGATTTTCC	TGATAAAAAT	900
CGCTTTAGAA	AATAATCTCC	TCAATTTTCAT	AGAGTTTGAA	AATGAGTGAG	ATTTTTTATT	960
TGCCCTTTGA	CAACTGAATA	GCCTAAAATG	GTACTTTCCT	CATTTGTGGA	GCAAATTTGA	1020
ATGGCTCGCC	ATGATAAGAG	CGATTTTAAA	ATCATCAATA	AAATAGAGCG	ATACTTTATA	1080
TGCCATGATA	CAAATGATAT	ACAATGATAC	TTCTGACCGT	TCAGCCTGCC	AACGTAAAAG	1140
AGCAGCAAGT	GAAATCTTTA	TGATGACTTC	ATCAGTCATG	CCACGTTGAA	TGTGTGAGTT	1200
TGTTAGATAA	ACGCAATTA	TCCTCAAAAAG	GTTCCCGGAA	CCTTTTGAGT	TCTACAGACG	1260
CATCACGTGG	AGTGTGTAAG	CTTGTTGCTA	AAAGCGTAAA	AACCTTGGA	CGAAAGGAAT	1320
AATAGACTTT	CTGCGAAAACA	AAAATATAAT	ACAATAAAAC	TATGAATGAT	GAAGCAAGTA	1380
AACAATTGAG	CGATAGCCGT	TTCAAGATCC	TTGTAGGTGT	TCAGCGCACG	ACTTTTGAAG	1440
AGATGTTAGC	TGTGTTAAAA	ACAGCTTATC	AACGTAACG	CGCAAAGGT	GGACGAAAAA	1500
GCAAATTAAG	CCTAGACGAT	CTCCTTATGG	TAACTATTCA	ATACATGCGA	GAATAGAGCA	1560
CTTATGAACA	AATTGCGGCT	GATTTTGGCA	TTCACGAAAG	CAACTTAATC	CGTCGGAGTC	1620
AATGGGTTGA	AGCAACTCTT	ATTCAAAATG	GTTTTACGAT	TTCAAATTCT	GCCTTAATTC	1680
TGTA AAAACA	GTA AAATTCG	AAGGATGTA	AGGTAAGAGT	TTTTTTCTTT	CTGAAAAAAT	1740
GGTATAATAG	CAATCAAAAC	TAGAAAATAA	AACGGAATTT	GGAACAGATT	TGTCTGTATC	1800
CTAGTAGAGT	GGTGATACTA	TGAAGATTAG	TAAGAGGCAC	TTATPAAATT	ATTCCATCTT	1860
GATCCCTAC	TTGCTTTTAT	CTATTTTGGG	CTTGATTGTG	GTCTATTCGA	CCACCAGTGC	1920
TATTTAAT	GAAGAAGCA	AGAGCGCCTT	GCAGTTGGTT	CGAAACCAAG	GAATCTTTTG	1980

GATTGTTAGT TTGATACTGA TTGCCTTAAT TTATAAATTG AGACTAGATT TTTTGAGAAA 2040
 TGAGCGACTA ATCATTTTAG TTATATTAAT AGAAATGCTT TTATTGTTCT TGGCTCGTTT 2100
 TATTGGTATT TCCGTAAACG GGGCATAACG TTGGATTTCG GTTGCAGGAA TAACTATTCA 2160
 GCCAGCTGAG TACTTAAAAA TCATTATTAT TTGGTATTTA GCTCACCGAT TCTCCAAACA 2220
 GCAAGAAGAA ATAGCTACTT ATGATTTTCA AGTTTGGACT CAAAATCAAT GGCTTCCCCG 2280
 TGCTTTTAAT GATTGGCGAT TCGTTCTCCT AGTTCGATT GGAAGTTTGG GAATTTTCCC 2340
 TGATTTAGGA AATGCGACTA TTTTAGTCTT GGTTCCTTG ATTATGTATA CAGTTAGTGG 2400
 AATCGCTTAT CGCTGGTTTT CAACCATTCT GGCCTCGTA TCTGCCGCTT CTGTCTTTGT 2460
 CTTGACCACT ATCAGCCTAA TCGGTGTTGA GACCTTTTCA AAAATCCAG TATTCGGCTA 2520
 TGTAGCCAAG CGCTTTAGTG CCTTTTTTAA TCCTTTTGCC GATCGTGCTG ATGCAGGTCA 2580
 CCAGTTAGCT AATTCCTATT TTGCCATGGT CAATGGCGGT TGGTTTGGTC TAGGTCTTGG 2640
 AAACTCGATT GAAAAACGAG GTTATTTGCC AGAAGCTCAT ACAGACTTTG TCTTTCTAT 2700
 CGTGATTGAA GAATTTGGCT TTGTTGGTGC CAGTCTTATT TTAGCTCTCT TGTTTTTCAT 2760
 GATTTTGGCG ATTATCTTGG TCGGTATCCG AGCGGAGAAT CCTTCAATG CCATGGTTGC 2820
 ACTCGGTGTC GGAGGGATGA TGTGTTTCA GGTATTTGTC AATATCGGAG GGATTTGGG 2880
 CTTGATTCCA TCTACAGGAG TGACTTTCCC CTCTTTATCC CAGGGTGGAA ATAGTCTTCT 2940
 AGTCTTATCA GTGGCAGTAG CCTTGTCTT AAATATTGAT GCCAGTGAAA AACGCGCTAA 3000
 ATGTACCBA GAATTTGAAA ATCAACCAAT GAACCTCTG TTGAAGTAGG ATAAAGAAAG 3060
 GATAGTTTAT GTCTCTTCAA AAATTAGAAA ATTATAGTAA TAAAAGTGTT GTGCAAGAAG 3120
 AAGTCTTGAT TCTAACAGAA TTAAGGGAAG ATATTACTAA AAATATGCTT GCCCAGAGA 3180
 CCTTTGAAAA AATAATACAG TTGAAAGAAT TATCAACGCA GGAAGATTAT CAAGGTCTAA 3240
 ACCGTCTAGT GACTAGCTTA TCAAATGATG AAATGGTCTA TATTTACGC TATTTCTCTA 3300
 TCTTGCCTCT TTTGATTAAT ATTTAGAGG ATGTGGATTT AGCTTATGAA ATCAATCATC 3360
 AAAATAATAT TGATCAGGAC TATTTAGGTA AATTATCTAC AACGATTAAA TTGGTAGCAG 3420
 AAAAGGAAAA TGCCGTTGAG ATCCTAGAAC ACTTGAATGT TGTCCCTGTT TTGACAGCCC 3480
 ATCCAACACA AGTGCAACGC AAAAGTATGT TGGATTTAAC AAATCATATT CATAGTCTTT 3540
 TGCCTAAATA CCGTGATGTT AAGTTGGGGT TGATCAATAA AGATAAATGG TACAATGATT 3600
 TGCGTCGTTA CATCGAAATT ATCATGCAGA CAGACATGAT TCGTGAGAAA AAAT'AAAAG 3660
 TGAATAACGA AATCACGAAT GCTATGGAAT ATTATAACAG CTCCTTTTGG AAAGCTGTAC 3720

776

CTCATTTGAC	GACGGAGTAT	AAGCGCTTAG	CGCAAGCGCA	TGGTCTGAAT	TTAAAAACAGG	3780
CTAAACCAAT	CACCATGGGT	ATGTGGATAG	GTGGTGACCG	TGATGGAAAT	CCATTTGTTA	3840
CAGCAAAGAC	CTTGAAGCAG	TCTGCACTCA	CTCAGTGTGA	AGTCATCATG	AACTACTATG	3900
ATAAAAAGAT	TTACCAACTT	TATCGTGAAT	TTTCTCTTTC	AACTAGCATT	GTCAACGTCA	3960
GCAAGCAAGT	CAGAGAAATG	GCTCGTCAAT	CCAAGGATAA	CTCGATTTAC	CGCGAAAAAG	4020
AGCTTTACCG	TCGTGCCTTG	TTTGATATTC	AATCAAAAAT	TCAGGCAACT	AAAACCTATC	4080
TGATTGAGGA	TGAAGAAGTT	GGGACTCGTT	ATGAAACCGC	CAATGATTTT	TACAAGGATT	4140
TGATTGCCAT	TCGAGATTCT	CTACTAGAAA	ATAAGGGCGA	GTCCTTGATT	TCAGGTGATT	4200
TTGTGGAATT	ATTGCAGGCA	GTAGAGATAT	TTGGTTTTTA	CTTAGCATCA	ATTGATATGC	4260
GACAAGACTC	TAGCGTCTAT	GAAGCCTGTG	TGGCAGAACT	CTTGAAATCA	GCAGGAATTC	4320
ATTCTCGTTA	TAGCGAGTTG	AGCGAAGAAG	AAAAGTGTGA	CCTTCTCTTG	AAAGAATTAG	4380
AAGAAGATCC	CCGAATTCCT	TCTGCGACTC	ACGCAGAAAA	ATCAGAATTA	TTAGCAAAAAG	4440
AATTAGCTAT	TTTTAAGACG	GCTCGTGTTC	TGAAAGATAA	GTTGGGAGAT	GATGTCATCC	4500
GTCAGACCAT	CATTCACAT	GCAACCAGCC	TTTCTGATAT	GCTAGAATTA	GCTATCTCTGT	4560
TAAAAAGAAGT	AGGACTGGTG	GATACGAAA	GGGCGCGTGT	TCAGATTGTT	CCCCTTTTTG	4620
AAACAATTGA	AGACTTGGAT	CATTCAGAGG	AAACAATGAG	AAAATATCTT	TCTCTTAGCC	4680
TTGCCAAAA	ATGGATTGAC	TCACGAAATA	ACTACCAAGA	AATCATGCTT	GGCTACTCTG	4740
ACAGTAATA	AGATGGCGGT	TACTTGTCAT	CATGTTGGAC	CCTCTACAAG	GCTCAACAAC	4800
AATTGACTGC	TATTGGAGAT	GAATTTGGCG	TTAAGGTTAC	CTTCTCCAT	GGTCGTGGTG	4860
GTACTIONCG	TCGTGGTGGT	GGGCCAACCT	ATGAAGCCAT	TACATCTCAA	CCGCTCAAGT	4920
CTATCAAGGA	TCGTATCCGC	TTGACGGAGC	AGGGTGAAGT	AATTGGGAAT	AAATACGGTA	4980
ACAAAGACGC	CGCTTACTAT	AACCTGAAA	TGCTAGTATC	GGCAGCTATT	AACCGTATGA	5040
TTACTCAGAA	GAAGAGCGAT	ACCAATACCC	CAAATCGTTA	TGAAACCAT	ATGGATCAAG	5100
TAGTGGACCG	TAGTTACGAT	ATCTACCGTG	ATTTGGTCTT	TGGTAATGAG	CATTTCTATG	5160
ATTATTTCTT	CGAGTCAAGT	CCAATCAAGG	CTATTTCAAG	TTTTAATATT	GGTTCTCGTC	5220
CAGCCGCTCG	TAAGACTATT	ACTGAAATCG	GTGGTTTGCG	TGCCATCCCT	TGGGTATTCT	5280
CATGGTCACA	GAGTCGTGTT	ATGTTCCCTG	GATGGTACGG	GTTGGTTCA	AGCTTCAAGG	5340
AATTTATCAA	TAAAAATCCA	GAGAATATTG	CTATCTTACG	AGATATGTAC	CAAAATTGGC	5400
CTTCTTCCA	ATCGCTTCTT	TCAAATGTTG	ATATGGTTTT	GTCAAAATCA	AATATGAATA	5460
TTGCTTTTGA	ATATGCTAAA	CTTTGTGAAG	ACGAGCAAGT	TAAGGCCATC	TATGAGACTA	5520

777

TTTTAAATGA ATGGCAAGTT ACTAAGAACG TTATCTTGGC TATTGAAGGA CATGACGAAC 5580
 TCTTAGCTGA CAATCCATAT CTAAAAGCTA GTCTGGATTA CCGTATGCCT TACTTTAATA 5640
 TTCTCAACTA TATTCAGTTG GAGTTGATTA AACGCCAACG TCGTGGAGAA TTGTCCAGTG 5700
 ATCAAGAACG ATTGATTCAT ATCACCATCA ACGGAATTGC GACAGGATTC CGTAATTCAG 5760
 GTTGATAATT TTCAGAGTG AATGCTAAAA GTGAATATCA AAAAAATTCT AATAGACTAT 5820
 TGACAAGTAG TTTAAAAATG ATATAATTTA ACCATTCAGA AAAGTAATCA TACAACTTT 5880
 TTAGAGAGTC TGTGGTAGCT GAAAACAGAT AAGTGGCAAT GATGAAAATT GGGCTGAATG 5940
 CTATTTAGAA TTTGAAAATTA TAAAAATTCG GTAAGCACAC CTTACAGTGC ATCTCGTTAT 6000
 TGCGGAGACTG AGCGATAGGG AAATTCCTTA TAATTGAGGT GGTACCGCGC ATCGACGTCC 6060
 TCACACAAGT TTTTGTGTG AGGATTTTTT TGATGGAGGT TAGTATGGAA AGAAAACGAT 6120
 GCGCTCGCTT GTTTAGATAA GTGAAATATG TTAAAGGAAA TAAAAGGAG AACAGAATG 6180
 AAAAAATAAC GTTTAATTGG AATTATTGCT GCATTAGCAG TCTTAGTAGC AGGAAGCTTG 6240
 ATTTATCTTT CAATGAATAA ATCAGAAGCT CAGAATAATA AGGATGAGAA GAAAAAACC 6300
 AAGATTGGTG TGCTTCAATT TGTGAGCCAT CCATCCCTTG ATTTGATTTA TAAAGGGATC 6360
 CAAGATGGAC TTGCAGAAGA AGGATATAAA GATGATCAAG TTAAAATTGA TTTTATGAAC 6420
 TCAGAAGGTG ACCAAAGTAA GGTTCGACA ATGAGTAAAC AATTGGTTGC AAATGGGAAT 6480
 GACCTTGTGG TTGGTATCGC AACACCAGCA GCCCAAGGGT TGGCTAGTGC AACAAAAGAC 6540
 CTACCGGTTA TCATGGCCGC TATTACAGAC CCAATTGGTG CTAAGTTGGT TAAAGATTTG 6600
 AAAAAACCAG GTGGCAACGT TACAGGGGTA TCTGACCACA ATCCAGCTCA ACAACAAGTT 6660
 GAACTCATCA AGGCTCTGAC ACCGAATGTG AAAACAATCG GAGCTCTTTA CTCAAGTAGC 6720
 GAAGACAATT CAAAA 6735

(2) INFORMATION FOR SEQ ID NO: 105:

- (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 6516 base pairs
 (B) TYPE: nucleic acid
 (C) STRANDEDNESS: double
 (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 105:

CTAGAGGATC CCAGCAGGTA AATTGGCTTC AGCTGGCAAA AAAGTTGCC TC GTTGAACG 60
 CAGCAAGGCT ATGTACGGTG GAACTTGTAT CAACATTTGGT TGTATCCCAA CTAAAACCTT 120

778

GCTAGTTGCT	GCTGAAAAGG	ACTTGTCTTT	TGAAGAAGTC	ATTGCTACTA	AAAACACGAT	180
CACTGGTCGC	CTCAACGGTA	AAAACATATGC	GACTGTTGCT	GGTACAGGCG	TAGATATCTT	240
TGATGCGGAA	GCTCACTTCC	TTTCAAATAA	AGTCATCGAA	ATCCAAGCTG	GTGATGAAAA	300
GAAAGAAGT	ACTGCTGAAA	CAATCGTCAT	CAACACTGGT	GCTGTTTCAA	ACGTCTTGCC	360
AATCCCTGGA	CTTGCTACAA	GCAAAAACAT	CTTTGACTCA	ACAGGTATCC	AAAGCTTGGA	420
CAAATTACCT	GAAAAACTTG	GAATCCTTGG	TGGCGGAAAT	ATCGGTCTTG	AATTTGCCGG	480
CCTTTACAAC	AAACTTGGA	GCAAGGTCAC	AGTCTTAGAT	GCCTTGGATA	CATTCCTACC	540
TCGTGCAGAA	CCTTCCATCG	CAGCTCTTGC	TAAACAATAC	ATGGAAGAAG	ATGGCATTGA	600
ATTGCTTCAA	AATATCCATA	CTACTGAAAT	CAAAAACGAT	GGTGACCAAG	TGCTTGTCGT	660
AACTGAAGAC	GAAACTTACC	GTTTCGACGC	CCTTCTCTAC	GCAACTGGAC	GCAAACCAA	720
TGTAGAACCA	CTTCAACTTG	AAAATACAGA	TATTGAACTA	ACTGAACGTG	GTGCTATTAA	780
AGTAGACAAA	CACTGTCAA	CAAACGTTCC	TGGTGTCTTT	GCAGTTGGAG	ATGTCAACGG	840
TGGCCTTCAA	TTFACTTACA	TTTCACTTGA	TGACTTCCGT	GTGTTTACA	GCTACCTTGC	900
TGGAGATGGC	AGCTATACAC	TTGAAGACCG	TCTCAATGTG	CCAAATACTA	TGTTTCATCAC	960
ACCTGCACTT	TCACAAGTTG	GTTTGACTGA	AAGCCAAGCA	GCTGATTTGA	AACTTCCATA	1020
CGCTGTTAAG	GAAATCCCCG	TTGCAGCAAT	GCCTCGTGGT	CACGTAAATG	GAGACCTTCG	1080
CGGTGCCTTC	AAAGCTGTTG	TCAATACTGA	AACAAAAGAA	ATTCTTGGAG	CAAGCATCTT	1140
CTCAGAAGGT	TCTCAAGAAA	TCATCAACAT	CATCACTGTT	GCTATGGACA	ACAAGATTCC	1200
TTTACTTAC	TTTCAAAAAC	AAATCTTAC	TCACCCAACC	TTGGCTGAGA	ACTTGAATGA	1260
CTTGTTTGCG	ATTTAAGTTG	AGATTTAATC	GTATCGAACA	GCCCTCTTTG	GGCTGTTTTT	1320
ACTTCTGCGG	AATCTCAAAT	CTGTCTTCT	CCTCTTTTAT	GATATAATAG	AAACATGAAC	1380
TTAAAAACTA	CTTTGGGCCT	TCTTGCTGGG	CGTTCTTCCC	ACTTCGTTTT	AAGCCGTCTT	1440
GGACGTGGAA	GTACGCTCCC	AGGAAAAGTC	GCCCTTCAAT	TTGATAAAGA	TATTTTACAA	1500
AACCTAGCTA	AGAACTACGA	GATTGTCGTT	GTCACTGGAA	CAAATGGAAA	AACCCTGACA	1560
ACTGCCCTCA	CTGTGCGCAT	TTTAAAAGAG	GTTTATGGTC	AAGTTCTAAC	CAACCCAAGC	1620
GGTGCCAACA	TGATTACAGG	GATTGCAACA	ACCTTCCTAA	CAGCCAAATC	TTCTAAAAC	1680
GGGAAAAATA	TTGCCGTCCT	CGAAATGAC	GAAGCCAGTC	TATCTCGTAT	CTGTGACTAT	1740
ATCCAGCCTA	GTCTTTTGT	CATTACTAAT	ATCTTCCGTG	ACCAGATGGA	CCGTTTCGGT	1800
GAAATCTATA	CTACCTATAA	CATGATATTG	GATGCCATTC	GGAAAGTTCC	AACTGCTACT	1860
GTTCTCCTTA	ACGGAGACAG	TCCACTTTTC	TACAAGCCAA	CTATTCCAAA	CCCTATAGAG	1920

TATTTTGGTT	TTGACTTGGA	AAAGGGACCA	GCCCAACTGG	CTCACTACAA	TACCGAAGGG	1980
ATFCTCTGTC	CTGACTGCCA	AGGCATCCTC	AAATATGAGC	ATAATACCTA	TGCAAACCTG	2040
GGTGCCTATA	TCTGTGAAGG	TTGTGGATGT	AAACGTCCTG	ATCTCGACTA	TCGTTTGACA	2100
AAACTGGTTG	AGTTGACCAA	CAATCGCTCT	CGCTTTGTCA	TAGACGGCCA	AGAATACGGT	2160
ATCCAAATCG	GCGGGCTCTA	TAATATCTAT	AACGCCCTAG	CTGCTGTGGC	CATCGCCCGT	2220
TTCCTAGGTG	CCGATTCGCA	ACTCATCAA	CAGGGATTTG	ACAAGAGCCG	TGCTGTCTTT	2280
GGACGCCAAG	AAACCTTTCA	TATCGGTGAC	AAGGAATGTA	CCCTTGCTTT	GATTAAAAAT	2340
CCAGTCGGTG	CAACCCAAGC	TATCGAAATG	ATCAAACCTAG	CACCTTATCC	ATTTAGCCTA	2400
TCTGTCCCTC	TTAATGCCAA	CTATGCAGAT	GGAATTGACA	CTAGCTGGAT	CTGGGATGCA	2460
GACTTTGAAC	AAATCACTGA	CATGGACATT	CCTGAAATCA	ACGCTGGCGG	TGTTTCGTCAT	2520
TCTGAAATCG	CTCGTCGCCT	CCGAGTGACT	GGCTATCCAG	CTGAGAAAAT	CACTGAAACG	2580
AGTAATCTGG	AGCAAGTTCT	CAAGACCATT	GAGAATCAAG	ACTGCAAGCA	TGCCTATATT	2640
CTGGCAACTT	ATACTGCCAT	GCTGGAATTT	CGTGAACCTG	TGGCTAGTCG	TCAGATTGTT	2700
AGAAAGGAGA	TGAACTAATG	GTTTATACTT	CACTTTCCTC	AAAAGATGGC	AATTACCCCT	2760
ATCAGCTCAA	CATTGCCAC	CTCTACGGAA	ATCTCATGAA	TACTACGGGG	ACAATGGAAA	2820
CATCCTCATG	CTCAAGTATG	TGGCTGAAAA	ACTGGGAGCC	CATGTGACCG	TTGACATCGT	2880
TTCTCTCCAT	GATGACTTTG	ATGAAAATCA	CTACGACATC	GCCTTTTTCG	GTGGTGGTCA	2940
AGACTTTGAA	CAAAGTATCA	TTGCAGACGA	CCTACCTGCT	AAAAAAGAGA	GCATTGACAA	3000
CTACATCCAA	AACGACGGTG	TAGTTCCTGG	TATCTGCGGT	GGTTTCCAAC	TATTGGGTCA	3060
ATATTATGTT	GAAGCTTCAG	GAAAACGTAT	CGAAGGGCTA	GGGGTCATGG	GACACTACAC	3120
GCTCAACCAG	ACCAATAACC	GTTTTATCGG	TGACATCAAG	ATTCACAATG	AAGATTCGA	3180
TGAAACCTAC	TATGGATTTG	AAAATCACCA	AGGTCGTACC	TTCTCTCTG	ATGACCAAAA	3240
ACCGCTGGGA	CAGGTTGTCT	ATGGAAATGG	AAACAACGAA	GAAAAGGTCG	GTGAAGGGGT	3300
TCATTATAAG	AATGTCTTTG	GTTCCCTACTT	CCACGGGCCT	ATCCTCTCTC	GTAATGCCAA	3360
TCTGGCTTAT	CGCCTAGTTA	CTACTGCCCT	CAAGAAGAAA	TATGGTCAGG	ACATCCAACT	3420
CCCTGCCTAT	GAGGACATTC	TCAGCCAAGA	AATCGCTGAA	GAGTACAGTG	ACGTCAAAAAG	3480
CAAGGCTGAC	TTTTCTTAAA	CAAAGGAAAA	TGATATCAAA	GAACCTCCGTT	ATCTTGTCGG	3540
AGTTTTTTGT	CTTTTCTTTT	ACCCTTCTCC	CTTGCATTTT	CTCTCATTTT	TTGCCAAAAT	3600
AGAGGGGTAG	AAAGAAGGTA	GCATATGTCT	AAATTACAAC	AAATCCTAAC	ATATCTTGAA	3660

780

TCAGAAAAAC	TAGACGTTCG	TGTCGTATCT	GACCCCGTCA	CAATCAATTA	CCTCACTGGT	3720
TTTTACAGTG	ATCCCCATGA	ACGCCAAATG	TTCCTCTTTG	TCCTAGCAGA	TCAGGAACCT	3780
CTCCTCTTTG	TCCCAGCTCT	TGAAGTAGAA	CGTGCAAGTA	GCACCGTTTC	CTTCCCAGTA	3840
GTGGGCTATG	TCGATTCTGA	AAATCCATGG	CAAAAAATCA	AACATGCTCT	TCCACAACCT	3900
GACTTCAAAC	GTGTCGCTGT	TGAGTTTGAC	AATCTCATCT	TGACCAAATA	CCATGGTTTG	3960
AAAACAGTTT	TTGAGACTGC	TGAGTTTGAC	AACCTCACTC	CTCGTATCCA	ACGCATGCGC	4020
CTCATCAAAT	CAGCTGATGA	AGTGCAAAAA	ATGATGGTTG	CAGGTCTTTA	TGCTGACAAG	4080
GCTGTTCATG	TTGGTTTTGA	CAATATTTCT	CTTGATAAGA	CTGAGACAGA	TATCATCGCA	4140
CAAATCGACT	TTGCCATGAA	ACGTGAAGGT	TATGAAATGA	GCTTTGATAC	CATGGTCTTG	4200
ACTGGTGATA	ATGTGCGGAA	TCCACACGGC	ATTCCAGCAG	CTAATAAGGT	TGAAAATGAT	4260
GCTCTTCTCC	TCTTTGACCT	GGGTGTTCTG	GTCAATGGCT	ATGCGTCAGA	TATGACTCGT	4320
ACAGTCGCTG	TCGGCAAACC	AGACCAATTC	AAGAAAGATA	TTTACAACCT	GACTCTTGAA	4380
GCCCAACAAG	CTGCTCTTGA	CTTTATCAAG	CCAGGTGTGA	CTGCTCATGA	AGTGGACCGC	4440
GCTGCCCGTG	AGGTCAATCGA	AAAAGCTGGT	TATGGTGAAGT	ACTTCAACCA	CCGTCTCGGG	4500
CATGGTATCG	GTATGGATGT	CCATGAATTC	CCATCTATCA	TGGAAGGAAA	CGACATGGTC	4560
ATCGAAGAAG	GCATGTGCTT	CTCTGTTGAA	CCAGGTATCT	ATATCCCTGG	TAAAGTCGGT	4620
GTTCGTATTG	AAGACTGCGG	TGTTGTTACC	AAGGATGGCT	TCAACCTCTT	TACAAGCACC	4680
AGCAAAGATT	TGCTTTATTT	TGATTAAACT	ATATAGCCCC	TATGCTTTCC	TTTCAAAATA	4740
TCTAGGGGCT	ATTTTATTGT	CATTTTCTG	CTATATGCT	AAAGAAATTG	GCTGCAATAA	4800
TCTAACCCTA	AGTGTCTGGA	ATGATAACGA	GGGTGCTCTC	CGCTTTTATC	AAAGACAAGG	4860
GATGAAACCC	CAAGAAACAA	CAATGGAAAT	GATAATTGAT	TAAGAAGTCA	TCTATCAAAA	4920
GATGTTAGAA	AAAGTTCAAT	TTCACTAGAA	AATGAGGAAA	ATCTCCCCAC	AATAAACCGC	4980
ATAGTATCAG	GTATTGTTGA	CTGACCCCAA	ACAGTTAGAC	AATTAATTTA	TCCGAAGGAT	5040
TTAGTTCTGT	ACTGCACAGG	ACTAAGTCCT	TTTAGTTTTA	CCTTAATTCG	TTTGTGTTG	5100
TAGTAATCAA	TATAGTCTAT	AATGACTTGT	TCCAATTGGT	TAAGTGATTT	AAATGTTTTC	5160
TCATAGCCAT	AAAACATTTT	GGATTTTAAA	ATGCCAAAGA	AAGATTCAT	CATACCGTTG	5220
TCTTGGCTGT	TTCCCTTGCG	TGACATAGAT	GCTTGAATTC	CCTTATTCTC	TAGGAACCGA	5280
TGATAAGAAT	CGTGTGGTA	TTGCCAGCCT	TGGTCACTAT	GGAGAATCGT	ATTCTCGTAG	5340
TGCTTCTCTT	TGAATGCCTG	TTCCAACATF	GTTTGTACTT	ATTCTAAATF	AGGCGAACAA	5400
GAAAGATTAA	AAGCAATAAT	TTCGCTGTTA	AAGCCATCTA	AAACTGGTGA	TAAGTAAAGC	5460

781

TTTGGAGTAC TTGCTGGAAT GGCAAATTCA GTCACATCTG TGTAGCACTT TTCCATTGTT 5520
 TTAGAGCCTT CAAATTGGGC TTGAATGAGA TTCTCTGCCT TCTTACCAAC GTCCTCTTTA 5580
 TGAGAAGAAT ATTTTCGTTT CTTTCGCATT TTAGCTTGTA AATTGAGTAC TTTTCATCAAG 5640
 CCTTGAAGTC TTTTATGATT TACCAGATAA CCACGATTTC TTAGTTCTAA ATGAACCCGG 5700
 CGATAAGCAT AATTTCCCTT GTGTTTCGATA AAGATGGATT GAATTTTCAGT TTTAAGCTCT 5760
 TGGTCTTTAT CTGTTTTGTC TAGCTGTTTC AAGTGATAGT AGTAGGTCCA ACGAGCTAGT 5820
 TTAATGGCTT CTAGAAGAAG ATCTAACGAA AACTCAGTCA TTAATTCCTG AACAAATTTCT 5880
 GTCTTTCTTC TTTCTCTTTT TCCTCCTTCA ATCGGAGTTC TCTTAACTTT TTTAGGATGG 5940
 CATTCTCCGC TCTCAGGTAC TCTCCCTCTT GTTTTCTCAA CAATAGTATA CCCGTTTTTC 6000
 CTGTATTGTG CTAGCCAGTT AAGAAGTATC GTACGACTTG GGAGACCGTA TTC AAGAGAA 6060
 ACTCTATCTT TAGTCCAGCC TTTCATGTCAG ACTTTATTAA CCCCAATTAT TCACCCCAA 6120
 TCTAAAACC ATCCAGAATC CTTGCCTTAG CTAGATCCT GGATGGTTTC TTTTTCACC 6180
 CAATGGGTGT TTTTACTAG AAAAAAGA GTTTCCCTT TATGGTATAA GTGTAGAAAA 6240
 AAACACAAA AGAAAGGAAA CTCACATGAA CAGTTTACCA AATCATCACT TCCAAAACAA 6300
 GTCTTTTAC CAACTATCTT TCGATGGAGG TCATTTAACC CAGTATGGTG GTCTTATCTT 6360
 TTTTCAGGAA CTTTTTCCC AGTTGAAACT AAAAGAGCGG ATTTCTAAGT ATTTAGTAAC 6420
 GAATGAmCAA CGCCGCTACT GTCGTTATTC GGATTCAGAT ATCCwTGTCC AGTTCCTCTT 6480
 TCAACTGTTA ACAGGTTATG GAACGGAATA TGCTTG 6516

(2) INFORMATION FOR SEQ ID NO: 106:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 14654 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 106:

TTTTCAACCC ATATCGTGGC TCCTGAATAC TACTTACTGA CAACTATGCT ATCAGAGACT 60
 TCTCTACTTG TTTTCTATAT CATTTCATC CATAGAAAAC AACTCATCCA CTTGGGACAT 120
 ATCTTTAGCT ATACTGTTTCG ATACTCTCTC TTTTCACTTT CCTTTGTAGC AATTTATTTT 180
 CTGATTAATT TCGTGTATCC TGTAGATATG GTCATTAATT TGCCATTTT GATTAATACT 240
 GGTTTGATTG TCTTGCTATC AGCTATCTCT TATATTAGTC TACTTGTCTT CACAAAAGAT 300

782

AGCATTCTCT	ATGAATTTT	AAACCATGTC	CTAGCCTTAA	AAAATAAAAT	TAAAAAATCA	360
TAGGAGTTTA	AAATGAAACA	ACTAACCCTT	GAAGATGCCA	AACAAATTGA	ATTAGAAAT	420
TTGGATTATA	TTGATACTCT	CTGTAAAAAG	CACAATATCA	ACTATATTAT	TAAGTACGGT	480
ACTCTGATTG	GGCGGTTTCG	ACATGAGGGC	TTTATCCCTT	GGGACGACGA	TATTGATCTG	540
TCCATGCCTA	GAGAAGACTA	CCAACGATTT	ATTAACATTT	TTCAAAAGGA	AAAAAGCAAG	600
TATAAGCTCC	TATCCTTAGA	AACTGATAAG	AACTACTTTA	ACAACTTTAT	CAAGATAACC	660
GACAGTACGA	CTAAAATTAT	TGATACTCGA	AATACAAAAA	CCTATGAGTC	TGGTATCTTT	720
ATCGATATTT	TCCCTATAGA	TCGCTTTGAT	GATCCTAAGG	TCATTGATAC	TTGTTATAAA	780
CTGGAAAGCT	TCAAACCTGCT	GTCTTTCAGT	AAACATAAAA	ATATTGTCTA	TAAGGATAGC	840
CTTTTAAAAAG	ATTGGATACG	AACAGCCTTC	TGGTTACTCC	TTCGACCGGT	TTCTCCTCGT	900
TATTTTGCAA	ATAAAATCGA	GAAAGAAATT	CAAAAATATA	GTCGTGAAAA	TGGGCAATAT	960
ATGGCTTTTA	TCCCTTCAA	ATTTAAGGAA	AAGGAAGTCT	TCCCAAGTGG	TACCTTTGAT	1020
AAAACAATCG	ATTTACCCTT	TGAGAATTTA	AGCCTTCCTG	CACCTGAAAA	ATTTGATACT	1080
ATTTTGACAC	AATTTTATGG	AGATTATATG	ACCCTACCAC	CAGAAGAAAA	ACGCTTCTAC	1140
AGTCATGAAT	TTCACGCTTA	TAAATGGGAG	GATTAGGATG	CAATATTTAG	AAAAAAAAGA	1200
AATTAAGAA	ATTCAACTAG	CCCTGCTGGA	CTATATTGAT	GAGACTTGTA	AGAAACATGA	1260
TATTCCTTAT	TTTCTCAGTT	ATGGAACCAT	GCTTGGAGCC	ATCCGCCACA	AAGGTATGAT	1320
TCCTTGGGAT	GATGATATTG	ATATTTCCCT	TTATCGTGAG	GATTATGAGC	GTTTACTGAA	1380
GATTATTGAA	GAAGAAAATC	ACCCTCGCTA	CAAGGTCTCT	TCCTACGATA	CATCTTCTTG	1440
GTACTTCCAT	AATTTTCGCAT	CGATTTTGGG	CACCTTCTACT	GTTATAGAAG	ACCATGTTAA	1500
GTACAAGCGT	CATGATACCA	GCCTTTTCAT	CGATGTCTTC	CCAATTGATC	GATTTACAGA	1560
CTTGAGCATT	GTCGACAAGA	GCTATAAGTA	TGTGGCTCTT	CGTCAACTAG	CTTATATCAA	1620
AAAATCACGA	GCAGTTCACG	GTGATAGCAA	ACTAAAAGAT	TTTCTTAGAT	TATGTAGCTG	1680
GTACGCTCTC	CGATTTGTCA	ATCCTCGCTA	CTTTTACAAG	AAAATTGATC	AACTAGTCAA	1740
AAATGCTGTA	ACCAACTCTC	CTCAATATGA	AGGAGGAGTT	GGGATCGGTA	AGGAAGGGAT	1800
GAAAGAAATC	TTCCCAAGTT	ATACCTTTAA	AGAACTGATT	TTAACTGAGT	TTGAGGGCCG	1860
TATGTTGCCT	GTTCCCAAAA	AATATGACCA	ATTTTAAACC	CAGATGTATG	GCGATTATAT	1920
GACACCACCA	TCAAAAGAAA	TGCAAGAGTG	GTATAGTCAT	AGCATTAAAG	CTTATCGCAA	1980
AAACTGATTG	AGGGGGATTA	TACAACTAC	TAAGATAGAG	GTTATTCAA	AACATAATTT	2040
TAGTAGAAAA	TGAAATACAT	ATTCACCAA	TAAAACGCAT	CATATCAAGG	TTTTTGAAAA	2100

ACCTTGATAT GATGCGTTTT ATAATTTTAA AGACTTTTTT CTATAGTAGA TTGAAATAAG 2160
 ATGCGAACAA ATCAATTAGA AAATTCAAAT TAATTTATAG AAATATTTTA GTATTCCTGT 2220
 GTACTGTTCT AAATTCAGTC TGCTATATCT TATTTTCTA TTTAAATCGC TTCTGTAACA 2280
 AAGCTACGAC TTTCAAGTAC CTTAAGCATG GCATTAGCTG TATCTAGCGC TGTGAAGAGG 2340
 GGCACCCCGT GTTCAATGGC TGAACGACGA ATTTGCTCAC CATCTTCGTC AGCAGTTCGT 2400
 TTTGTTCCTA CTGTGTTAAT GATAGCTTGA ATTCTTCCTT TCGGTACAAA ACTTGGGATA 2460
 TCCTTATCGT CATCACCAAT CTTACCAACA GGTGGGCTT GCAAGCCATG ACTAGCAAAG 2520
 AAGGCTGCTG TCCCTTCTGT CGCAAGGATT CCATAACCAA TGTTTTGGAA ACGACGAGCC 2580
 AAGTCAAGG CTCTCTCTT GGCATCATCA GCGATGGTAA AGACGACATT ACCAAAAAGTT 2640
 GGCAAGTGTA GATAAGAAGC TTCAAAGGCT TTATAGAGAG CTTTTTCCAA AGTAGCATCA 2700
 GAACCCATAA CTTACCTGT TGACTTCATT TCAGGACCGA GCAAGCTGTC TACCTTAGCT 2760
 AGTTTGGTAA AGGAGAAGAC AGGTGCCTTG ATATGAACAC GGGTCTTTC AGGGTAAAGT 2820
 CCATTTGGT AGCCAAGTTC TGATAAACTT TGACCAAGAA TGAGTTGGT CGCTACTTGA 2880
 GCCATAGGAA TATTGGTTAC CTTAGATAGG AATGGAACAG TACGGCTGGC ACGTGGATTG 2940
 ACCTCAATAA CGTAGACTTT TTCATCCTTG ATAACAACT GGATGTTTAT CATTCGAAGG 3000
 CAGTGAAGAC CGATTGCTAA GCGTTTGGTG TAGTCTGCGA TGGTCTCCTG AACCTTTGTC 3060
 GACAAGGTTT GTGGTGGGTA AACAGCCATT GAGTCACCTG AGTGGACACC AGCACGTTTCG 3120
 ATATGCTCCA TGATACCAGG AATGAGTACA TTTTTACCAT CTGAAATGGC ATCAACTTCG 3180
 CACTCTTGCC CAACGATATA AGAGTCGACA AGAACTGGGT GGTCTGGACT AGCCTTAACA 3240
 GCAGTTCGCA TGTAAGAAGC AAGGTCTTCT TCGTTTTCAA CGATTTCCAT GGCACGTCCA 3300
 CCAAGTACAT AAGATGGGCG GACAAGAACT GGAAGCCAA TCTTGCAGC TGCAAGAGCT 3360
 GCTTCTTCTT CATTTGGTAGC CGTTTGTCTT GGTGGCTGTG GAATATCCAA TTCTTTGAGA 3420
 GCTTGCTCGA AGAGGTCACG GTCTTCGGCA CGATCTAGGT CAGCAACCTG TGTACCAAGG 3480
 ATGGTCACAC CTGCTTTTGC CAATGGCTCC GCAAGTTTGA TGGCTGTTTG ACCACCGAAC 3540
 TGAACGATAA CTCCCTTTGG TTGTTCGAAG TCAATGACGT TCATAACATC TTCGAATGTC 3600
 AATGGCTCAA AGTAAAGCTT ATCTGATACA GAGAAGTCTG TTGAAACGGT CTCTGGGTTT 3660
 GAGTTCATGA TGATAGCTT ATAACCAGCT GCCTGGATAG CCTTAACAGA GTGAACGGTT 3720
 GCGTAGTCAA ACTCAACCC TTGACCGATA CGGATGGAC CTGAACCTAG GACAAGTACA 3780
 GATTCTTTAT CAGATCTGAT AGATTCATTT TCCCAACCAT AGGTGGAATA GAAATATGGC 3840

GTTTCGGAGT	CGAACTCTGC	CGCACAAAGTG	TCTACCATCT	TATAAACTGG	AACAATCTTG	3900
TTTTCCAAGC	GAAGTTGGCG	AACTTTATCA	TCAGTCGTTC	CCCAGAGTTC	AGCAATCTTA	3960
CGGTCTGAAA	AACCATTAAG	TTTGGCTGTT	TTCAAACTT	CTAAATCTTG	TGGATGAGCA	4020
CCCAATCTCT	GCTCAATTC	AAAGATATGC	AAGAGTTTAT	CAAGATAGAA	GATATCAATT	4080
TTTGTAAGCT	CTGCAATTC	TTCAGGTGTG	TAGCCACGAC	GAATGGCTTC	TGATACGTAG	4140
AAGAGACGGT	CATCTTGGGC	TTTGACAACC	TTTTCAATCA	AGGCATCATC	AGAACTGCT	4200
GCAAGTTCAG	GTATTTCAAT	GTGGTGCACC	CCAAATTCOA	GGGAGCGGCA	GGCCTTGAGA	4260
AGAGATTCCT	CGATGTTACG	ACCGATTGCC	ATGACTTCTC	CAGTCGCCTT	CATTTGTGTA	4320
CCGAGACGGC	GTTCAACCCTT	TTCAAACCTG	TCAAATGGGA	AACGTGGAAT	CTTAGCAACT	4380
ACGTAGTCAA	GGCTGGTTC	AAACATGGCA	TAGGTTGAAC	CTGTAACCTG	GTTTATAACC	4440
TCATCCAAGG	TCAAACCTAC	TGCAATCTTG	GCAGCCAAC	TAGCAATCGG	ATATCCTGTC	4500
GCTTTAGAAG	CAAGGGCTGA	CGAACGTGAT	ACACGAGGGT	TTACTTCGAT	AACATAATAC	4560
TTGAAGCTGT	TAGGATCAAG	AGCTAGCTGA	ACATTACATC	CACCTTCAAT	CTTGAGGGCA	4620
CGAATAATGC	TCAAGCTCGC	ATCACGAAGC	ATTTGGTTTT	CATAGTCTGA	CATGGTTTTGC	4680
GCAGGGGCAA	ATACAATGGA	ATCCCCTGTG	TGAATCCCAA	CTGGGTCAA	GTTTTCCATG	4740
TTACAAACAA	CCAAGGCATT	GTCAGCTGAG	TCACGCATCA	CTTCGTATTC	AATTTCCCTG	4800
AAACCCGCAA	TCGAACGCTC	AATCAAACAT	TGGGTAACAG	GTGACAATTT	CAAACCATTT	4860
TCAGTGATTT	CACGCAATTC	TTTCTCGTTG	GCACACATAC	CACCACCAGT	ACCACCAAGG	4920
GTAAAGGCTG	GACGAACGAT	GA CTGGGTAG	CCAATTGTCG	CTGCAAAGGC	AACTGCTTCT	4980
TCTACTGTGT	TAACAATTC	AGATTCTGGA	ATGGGTGTT	CAAGCTCTTC	CATCAATGT	5040
TTAAAGAGGT	CACGGTCCTC	CGCTTGGTCA	ATGGCAGATA	ATTTGGTACC	CAGAAGTTCA	5100
ACGCCAAGCT	CGTCTAGGAT	ACCATTTTFA	GATAATTTCCA	TGGCCATGTT	GAGACCTGTC	5160
TGACCACCGA	GTGTTGGTAG	CAAGGCATCT	GGACCTTCCT	TACGAAGAAT	ACGTGTCACA	5220
AACTCAAGTG	TAATCGGTTC	AATGTAAACC	TTGTCAGCAA	TTTCCTTGTC	CGTCATGATG	5280
GTTGCAGGAT	TTGAGTTAAC	CAAACAACC	TCATAACCTT	CCTCTTTCAA	CGACAAGCAA	5340
GCCTGAGTCC	CAGCGTAGTC	AAACTCAGCA	GCCTGACCAA	TAATAATCGG	ACCAGAACCA	5400
ATCACCATAA	TTTTTTGAAT	ATCAGTACGT	TTAGGCATAT	ATAAGATATT	AAGGGTGTCA	5460
AGCGGACAAA	GCTAAAATAG	GAGTTATGAC	GAAGAAGTGT	CAGTTCTAGG	AATAACTATC	5520
TTTTTAGCAC	CGTCCGTAGC	CCGTATTCAG	TTCAGCAAAT	ACGGAGCACC	CTTCTCCTTT	5580
CTATTCGTCG	CCTCTCAGGG	CGACATTAFA	TAAGATACAA	AGGACGAATA	GAAAGCGATT	5640

785

GAATTTTAGG	AAATCAAGGA	AGGATGACA	ATCCAAGTTG	GTTTCTCTAC	ATTCTGAGCT	5700
TTCCGTCCGT	GTTCAGTTAC	ATAAAATTCTC	CGACGAGCTT	TTACTCGTTC	TTAGTTTGAT	5760
TGTTTAAAAA	CTTCCATCAT	CTCGATAAAC	TCGTCAAATA	GGTAGCTAGC	GTCGTGTGGC	5820
CCAGGAGCTG	CATCTGGGTG	GTATTGAACA	GAGAAAGCAG	GTTGGTATCT	GTGGCGCACA	5880
CCTTCCACTG	ACTTGTCAAT	GATTTCTTCG	TGGGTAATAA	TCAAGTGCTC	TGGCAAATCC	5940
TCGCGGCTGA	CTGCATAACC	ATGGTCTTGG	CTGGTGAAGT	CTACTCGTCC	TGTTGCGATT	6000
TCACGTACCG	CATGTTGAA	TCCACGGTGG	CCAACTTCA	TCTTATAGGT	CTTAGCCCCG	6060
TTTGCCATTG	CAAAGAGTTG	GTGTCCATA	CAAATACCAA	AGATGGAAAT	TTTTCTTGT	6120
ACACCGCGAA	TCATGTCGAG	TGCTTGTGGA	ACGTCTTCTG	GGTTACCTGG	ACCATTTGAC	6180
AACATAACTC	CGTCAGGATT	GAGATGGAGA	ATTTCTTCAG	CCGTTGTCGA	ATAAGGAACA	6240
ACTGTCACGT	TACAGTTGCG	TTTAGAAAGT	TCACGTAGGA	TTGAGTGCTT	GAGACCAAAG	6300
TCCACTAGCA	CCACGCTCAA	ACCAACTCCT	GGAGCTGGAT	AAGACGTTTT	AGTAGAAACC	6360
TGTTTGATAT	TGCTGTTCGG	TAAAAGTGT	GCTTGGAGCT	GGTCCGTAC	ATGGTCCATA	6420
CTGTCCCCAA	CATGGGTCAA	GGTGCACGC	ATAGTACCAT	GCTTACGGAT	AATCTTGGTA	6480
AGAGCACGCG	TATCAATTCC	TGAAATCCCT	GGAATTTTCT	TGGCTTCAA	AAATTCATCC	6540
AAGGTCATTT	GGTGCGCCA	GTTGCTAGCT	CTACGCGCTT	CTTCAAAAAC	AACGACTCCC	6600
TTACAAGTTG	GAATAATGGA	TTCATAATCA	TCACGATTAA	TACCATAAT	TCCTACCAAA	6660
GGATAAGTAA	AGGTCAAGAT	TTGTCCATTA	TAAGACTGGT	CTGTAATGGA	TTCTTGGTAG	6720
CCGGTCATCC	CTGTATTAAA	GACGATTTTCG	CCTGTTACAT	CAATATCTGC	TCCGAAGGCC	6780
TTGCCTTCAA	AAACTGTGCC	ATCTTCTAAT	ACTAGAATTC	TTTTTGTCAT	ATTTTCACCT	6840
CTCGTGGACG	CTCACTGGCG	TCTTTTAACG	TCTTGTGTTT	TAGTTGGCGT	TTCTACTCGC	6900
TAGTACGGAT	TCTAAGATTG	CCATTCGAAC	AAAGACACCA	TTGGTCATTT	GTTGGACAAT	6960
CCGTGATTTT	GGTGCTTCAA	CCAAGTGGTC	TGCTATTTTCT	ACATCACGAT	TGATTGGAGC	7020
TGGGTGCATG	AGGATTGCTG	TTTCTTCAA	ACGATCGTAA	CGTTCTTGAG	TCAAGCCATG	7080
TTGGGCATGG	TAGTCTTCTT	TTGAAAATAC	AGCTCCACTA	TCATGGCGTT	CGTGTGTCAC	7140
ACGGAGAAAC	ATCATGACAT	CAACCTGATC	AATGATTTCA	TCAATGGTTA	CAAAGTGTCC	7200
ATAGTCTGCA	AACTCTTGAC	TTCTCCATTC	CTCAGGTCCA	GCGAAAAGA	GTTGAGCTCC	7260
CAAGCGTTTC	AAAATCTGCA	TATTGGATTT	GGCAACGCGT	GAGTGGTCCA	AGTCACCTGC	7320
AATAGCAACT	TAAAGACCCT	CAAAGTGGCC	AAATCCTCA	TAAATGGTCA	TCAAATCAAG	7380

786

CAAGCTCTGG	CTAGGGTGT	GGCCGAACC	ATCTCCACCA	TTGATGATGG	AAGTCGTAAT	7440
CGTTGGACTA	GCAATCAATT	CTCTATAGTA	GTCGACCTCT	GGATGGCGAA	TCACACAGAC	7500
ATCCACTCCT	AAAGCAGACA	GAGTCAAAAT	GGTGTCAATA	AGTGTCTCAC	CCTTATTAAC	7560
CGAGCTAGTC	TTACATCAA	AGTCAAGTCG	TTCCAATCCA	AGTTTAATCT	CTGCGACTTC	7620
AAAGGACTTA	TGTGTCCGTG	TAGAATCCTC	AAAGAAGAGA	TTGGAAACAA	TCGGATGGTC	7680
TTCATAGGGA	AGCTGGGCTC	CATTTTTAAA	CTCAATTCCT	CGCTTGATCA	ATTTTCATTAC	7740
TTGATCGACA	GTGAGGTCTT	CCATGGACAC	CACATGGTTC	AATGCTTGTT	GATTTTCTGA	7800
CATGGCTACT	CCTTTAACTT	TCTAAGCTTC	TTCAAGTAATC	AGAACTCTGT	CTTGGTCATC	7860
AAGTTCTGTC	ATCTCTACGA	TGATTTCTTC	AGAACGACTG	GTTGGGATAT	TTTTTCCAAC	7920
GTAATCTGGA	CGGATTGGCA	ATTCTCTATG	TCCACGATCG	ACTAGAACTG	CTAAACTCAC	7980
ACGCGCAGGA	CGACCATGAC	CGACAATATT	ATCAATAGCA	GCACGGATGG	TACGACCTGT	8040
ATAGAGCACA	TCATCCACCA	AGATAACTTC	GCGGTCTGTC	ACATCGACAG	AAACCAAAGA	8100
AGTATCTTCT	CCACTTTTAA	CATCATCAG	GAAAGGTTTA	GTATCCAATT	CCACAACAGG	8160
AACTGAAAAG	TTTTCTAACT	GCTTCAAACG	TTCTTGATT	CGGTGGGCAA	TAAAGACACC	8220
ACGAGTTTTA	ATACCAGCCA	AGACGATCTT	ATTCAAATCT	TTGTTGCGTT	CGATAATCTC	8280
ATAAGTAATA	CGCGTAATCG	CTCGTTTGAC	GGTCAATTCG	TCTACAACCT	CTTTTGTTTT	8340
CATGACAAAC	CTCCAAAAG	AAAAGTCTCC	TFAACAAGG	AGACTTGAAA	TTTATAGCCA	8400
AGCGAGCCCT	ACTGCACACA	GTATAGACTT	CACCCTTCTA	CTTTATCGCG	CTCCTTGCCCT	8460
GCCTCACGGG	ACAGGTTTAA	AGGAATATTT	AGTTATCATT	TACTATAGCA	CAAAGCATGC	8520
TTAAAATCAA	GCAAAAAGTT	TCAATGTAGC	ATCTTACAAA	TTGCTAAAAT	CATATAATTG	8580
TGGGTACTGG	TCACACTCTG	GATTTTTTGG	ATGGCAAATG	GCTCTTCCAA	AATAAATCAT	8640
GGCCTGATGG	GCAGCTAACC	ACTGCTCAGG	CGGCAAGATA	TCCATGACCC	GCTTTTCCAC	8700
CTCAAGTGGC	GTCGCTGATT	TTTTGACAAT	ATCGTGGTGT	TTGCAAATAC	GCTCCACAIG	8760
AGTATCCACT	GCAAAGGCTG	GAATTCCAAA	TCCTACACTC	ATGACAACAT	TGGCTGTCTT	8820
GCGACCAACA	CCTGCCAAAC	TCTCCAATTC	TTACCGTGTC	TGAGGGACTT	GACCATCAAA	8880
ATCGTCTAGT	AACTGTTGGG	CACATTTTTT	AAGGAATTTA	GCTTTATTCC	GATACAATCC	8940
CAAGCGAGAA	ATATGTGAAG	CAATCTCACT	CTCTGTCGCT	ACAGACATAG	CTTGGGGTGT	9000
TGGAAAGGCA	ACAAAGAGAC	CTGGTGTGGC	CTTATTTACC	GCTGCATCTG	TCGTCTGGGC	9060
TGATAACATG	ACCGCAACCA	GGAGTTCAAA	ATGATTTGGTA	AAATCAAGAC	TAGGCTTGGC	9120
ATCTGGGAAG	AGGGCAATGA	TTTCTTCTAG	CACCTTTCGT	GCTCGTTTTT	TTGACAAGAC	9180

CATTATTCAT	CTCCGTCAA	TAGTCCTTGT	AAGCCAGCAA	AAGGACTGTT	TTCTTCTTTC	9240
TTTACTGCCT	TTTGGACTTG	GTATTCCTCC	TCTGTCATGA	TTTGCCAGTC	ATTTCCCTGAG	9300
ATAAATCCTT	GACCAGCTTC	TTCTTCAGCC	GTCAAGACCT	TGATAGGAAT	GTTTAGCAGG	9360
ATATTGTCTG	ATACACTCTC	AGCAAGGTCA	AGCTCCCCAT	TTTCGATGGG	CAAGACCAAG	9420
TCATCATCTA	AAACTTCTTG	ATCTAGCTGG	TTAGTTGCGC	CTTCCATGAA	AACTTCCGTG	9480
ACTGGATAAG	ATTCAACTAA	CTCAACTGGC	TCCATACTGC	GACTCGACGC	AAGAACAATG	9540
GTATAAGATA	GTTGATAATC	TAAGAAATAC	ATACGGTCTT	CATATTGTAC	TTTCCCAACT	9600
GCAAGGATAT	CTTTTACATC	TAAAATTTCT	TGATTACGTG	CACGCAGGTC	ATCAACTAAA	9660
TCTAACGTTT	GTTCAAAGTT	CAAACCTTCA	GACTGCTTAC	GAATTTCTTG	AATATTTAAT	9720
TTCATACTTC	CTCCATAAAG	ATTTACTCTC	TTGATTATAC	CATGAAAAGG	CTACAAATCA	9780
GCACACCAA	CTTTGTAATT	AAAATTCAAA	ATTTAACAAT	ATTTACTATG	ATAGTTTTAT	9840
TTTTTAGTGC	TATACTATAG	GGAAAGAGTA	CATCAGATCA	AGGAGGATGC	TCACATGGAA	9900
GACAAGAAAC	TCATTCAACT	CCTATCCAAG	TTAAATAAAA	GCTACCAAAA	CTGTAAACAG	9960
GGTACGGCAG	ATGATATTCG	ACTACAAGAG	CTGCTAAACA	CTACTATGCA	AGAGCTCAA	10020
AAAACGGAAC	AGTTGAACAA	CAGTATCTTA	ATTGATCTTG	AGAAATTTTA	CCAACCTACC	10080
AGTCTTCTGA	TTGGACTGGG	TAGCCTAAAA	CTAAACGATC	AAGCACGCAC	TGCTTGCGGA	10140
AACTATGATA	AATTCATTAA	CGATCATGTC	AAACACGTAC	TAAGTCTCTA	TGGACCTGTT	10200
TTTGAATTTT	AGAGCATAGA	ATTTCCAGTT	TTCTGTTGAC	AAAATTTTCT	TAAAGGTATA	10260
ATATAAAGAT	ACTAATACTC	GGAGGTAAGG	GAGACATGAA	CAACTAAGTC	TATCAAATAA	10320
AGAACCTTTA	TTTAGTAGAT	CTTGTTTTTG	TCTCTTTTTG	TGTGCTCTTT	TATGCTCTTT	10380
TTCTGGCATG	TTAATAGAGT	TTTTTTGACA	TAGACTTTGG	GCTCTACTAG	GTAAGTAGA	10440
GCTTTTTGTT	ATGCACTATG	AACATTCTAG	AAAGGGAAAT	CATATGATAA	AAATCAATCA	10500
TCTAACCATC	ACACAAAACA	AAGATTTACG	AGATCTTGTA	TCTGACCTAA	CCATGACCAT	10560
CCAAGACGGG	GAAAAGGTTG	CTATTATTGG	TGAAGAAGGA	AATGGCAAAT	CAACCTTACT	10620
TAAAATTTTA	ATGGGGGAAG	CTTTGTCTGA	TTTCACTATC	AAGGGAAACA	TCCAATCTGA	10680
CTATCAGTCA	CTGGCCTACA	TTCCCTCAAAA	AGTCCCTGAG	GACCTAAAAA	AGAAAACTTT	10740
ACACGACTAC	TTCTTTTTAG	ATTCTATTGA	TTTAGACTAC	AGTATCCTCT	ATCGTTTGGC	10800
GGAGGAATTG	CATTTTGATA	GCAATCGTTT	CGCAAGTGAC	CAAGAGATTG	GCAATCTATC	10860
AGGGGGCGAA	GCTTTGAAAA	TTCAGCTTAT	CCATGAGTTA	GCCAAACCCT	TTGAGATTCT	10920

ATTTTTAGAT GAACCTTCAA ATGACCTAGA CCTTGAGACA GTTGATTGGC TAAAAGGCCA 10980
 GATTCAAAAAG ACCAGGCAAA CCGTTATTTT CATTTCCCAT GATGAAGACT TTCTTTCTGA 11040
 AACGGCAGAC ACTATTGTTC ACTTGCGACT GGTCAAACAC CGTAAAGAAG CGGAAACGCT 11100
 AGTAGAGCAT TTAGACTATG ATAGCTATAG TGAGCAGAGA AAGGCTAATT TTGCCAAACA 11160
 AAGTCAGCAA GCTGCTAACA ACCAAAGAGC CTACGATAAA ACCATGGAAA AACATCGGAG 11220
 AGTTAAGCAA AATGTAGAAA CTGCGCTPCG AGCTACCAAA GATAGTACTG CCGGTGCGCT 11280
 ATTGCGTAAA AAGATGAAAA CTGTCCTCTC ACAAGAAAAA CGCTACGAAA AGGCAGCTCA 11340
 GTCCATGACT CAAAAGCCAC TTGAAGAGGA ACAAATCCAA CTTTTCTTTT CAGACATCCA 11400
 ACCATTACCA GCTTCTAAAG TCTTAGTCCA ACTGGAAAAA GAAAATTTGT CCATTGACGA 11460
 CCGAGTTTTC GTTCAAAAAC TACAACCTAAC TGTCCGTGGC CAAGAAAAAA TCGGTATTAT 11520
 CGGGCCAAAT GGTGTTGGGA AATCAACTCT GTTAGCCAAG TTACAGAGAC TTCTGAATGA 11580
 TAAAAGAGAG ATTTCACTTG GTTTTATGCC ACAAGATTAC CACAAAAAAC TGCAATTGGA 11640
 TTTATCCCA ATAGCCTATC TCAGTAAAAC TGGGGAAAAA GAGGAACCTAC AGAAAATCCA 11700
 ATCTCACCTA GCTAGTCTCA ATTTCAGTTA TCCAGAAATG CAGCATCAA TTCTGCTCCTT 11760
 ATCTGGCGGA CAACAGGGAA AACTCCTGCT TTTGGATTTA GTCCTGCGCA AACCAAACCTT 11820
 TCTCCTGCTG GATGAACCCA CACGAACTT TTCTCCCACT TCTCAACCCC AAATCAGAAA 11880
 ACTCTTTGCT ACCTATCCAG GCGGTCTCAT CACTGTTTCG CATGACCGTC GTTCTTTAAA 11940
 AGAAGTCTGC TCGATCATCT ATCGCATGAC AGAACACGGT TTGAAGCTAG TTAATTTAGA 12000
 AGATTTATAA ATTTGCAACA TAGCAAAAAT CCAGAGACGA CCTCTGGATT CTTTTACATC 12060
 TGTTTTAAAC GTTCAATCCG TTCTGAGATA GGTGGGTGGG TATAAAAGAG TTTTTGGAAC 12120
 CCCCCACCTT TCTTAGGATC ATTGATATAA AGGGCACTGC TAGCATCATC GACGTGGCGA 12180
 CTCATAGGTT TGCTATFGTC CAACTTATCT AGGGCATTTA TCATTCCCTG GGGATTGCGA 12240
 GTCAGTCTGA CACTAGATGC ATCTGCCAGA AATTCCCTCT GACGAGAAAT AGCGAGCTGA 12300
 ACCAAGGTTG CAGCGAGAGG TGCCAGTACA ATAGCTAGTA GGGAAACCAC TAGCATAATG 12360
 ATTTCAAGAC CATTTCCATC TCGGTCATCA TCACTTCGTC TGCGACCTGC TCCACCCAC 12420
 CACATCATA GACCTGCCAT ACTAGAAAGC ATGGTGATAG CACTAGCAAG GGCAACTGCA 12480
 ATAGTCGAAA TACGGATATC ATAATTACGA ATATGACTGA CTTTATGTCC CATAACAGCT 12540
 TCTAGTTCTT CACGATTCAT GATAGCTAGT AGACCTGAAG TCGCAGCAAC AGCCGCATTT 12600
 TGAGGATTAG AACCTGTCGC AAAGGCATTT AAGGCTGGAT CATCAATGAT GAAAACACGG 12660
 GGCATAGGAA TCTGAGCGAC CAGAGCCATA TCTTCCACTA CATGGTAGAG GTCTGGTGCC 12720

GTTTGCTCAT	CCACCTCAG	CGCTCCATTC	ATGGACATGA	CAATCTCTGT	CGATTGAAAA	12780
ATCATAGACA	AAGCGTAGAT	AAAGCCGATA	ATCAGTGCAA	TAACCAAACC	ACCAAGTCCA	12840
GATCTTATAA	AGAGATAACC	AACCGCATAA	CCAACAAGAG	CTAAGAGTAG	GAAAAATACC	12900
AGCAACAAAA	TCCAGGTTTT	TCGTTTATTG	CTTGCAATTT	GATCAAACAA	CATCTTAGTC	12960
ACCTAAACCG	CTAAATCAA	CTTTAGGAAC	CGACTTTTCC	TCTTCAGGTG	TTTGAAGGAA	13020
ATCTGCCGCT	TTAAATCCAA	ACATTCCAGC	GATAATATTG	CTCGGAAAG	TTTCTAATTT	13080
TACATTGTAG	TTGCTGACAA	CACTGTTATA	GAGTTGACGA	GAGTAAGAAA	TTTTATTTTC	13140
TGTGTTTGTC	AACTCCTCTT	GCAATTTAAC	AAAGTTAGCA	CTAGCTTTCA	AATCTGGATA	13200
GCTTCTGCA	ACTGCAAAAA	TACCTGAAAC	CTGACGAGTG	AGGGCATCAC	TGGCTTTCAT	13260
AGCTTCTGCT	GGTGAAGTCG	CTGCCGCCAC	TTGGTTACGT	AGTTCTGCCA	CCTTTTCAAG	13320
GGTAGAACCT	TCATATTTGG	CATAACCTTT	TACAGTCTCA	ATCAAGTTTG	GCAAGAGGTC	13380
ATTGCGACGT	TTCAACTGAA	CATCAATCTG	ACTCCAAGCC	TCCTFGGTTT	GCATACGATT	13440
TTTAACCAAA	CCGTATAGC	TAACAATCAC	AAAAATAACA	ATAAGAGCGA	TAACTCCAAG	13500
AATAATCCAA	GTCATAATAT	AAGTCCTTTC	TGCTTTTAGA	TTAGTACCAG	TATATCAAAT	13560
TTTCTATGAT	TGTGGTAAAA	TAAGATGATA	CTAAGAAGG	AAATAACTAT	GAAACCAAAA	13620
ACATTTTACA	ACTTGCTTGC	CGAGCAGAAT	CTTCCACTTT	CGGACCAGCA	AAAAGAACAA	13680
TTTGAACGTT	ATTTTGAGCT	CTTGGTCGAG	TGGAATGAGA	AGATTAATTT	GACGGCGATT	13740
ACGGACAAGG	AAGAAGTTTA	TCTCAAACAT	TTTTACGATT	CGATTGCACC	CATTCTTCAA	13800
GGTTTGATTC	CCAATGAAAC	TATCAAACCT	CTTGATATCG	GGGCTGGGGC	AGGATTTCCCT	13860
AGTCTACCAA	TGAAAATTCT	CTATCCGGAG	TTAGATGTGA	CCATTATTGA	TTCACTCAAT	13920
AAGCGCATCA	ACTTCCTACA	ACTCTTGGCT	CAAGAACTGG	ATTTGAACGG	AGTTCATTTTC	13980
TACCACGGAC	GTGCCGAAGA	TTTTGCCCAA	GACAAGAACT	TCCGTGCTCA	ATATGATTTT	14040
GTAACAGCTC	GTGCGGTTGC	CCGTATGCAG	GTCCTATCTG	AATTGACTAT	TCCCTACCTT	14100
AAGGTTGGTG	GCAAATATT	AGCACTCAAG	GCTAGCAATG	CGCCTGAGGA	ATTATTAGAA	14160
GCTAAGAATG	CCCTCAATCT	CCTTTTLAGT	AAGGTCGAAG	ACAATCTCAG	TACGCCCTAC	14220
CGAATAGAGA	TCCGCGCTAT	ATCACAGTGG	TAGAAAAGAA	AAAAGAAACA	CCAAATAAAT	14280
ATCCACGTAA	GGCTGGTATG	CCAAATAAAC	GCCCACTTTA	AATTTTTTAG	TAAACAAATG	14340
TTTACAAAAT	CAGCCTCGCT	CTTTTATTTT	TAGGCTCGGG	AAAAAATGAT	TTACAAAATC	14400
AGCCTCGCTC	TTTATTTTCT	AGGCTCGGGA	AAAAATGATT	TACAAAATCA	TTTTTTTCTG	14460

790

CTATACTATC CTAAGCAAAG GTTTTTAATG TCATCCCGTG AGGTGACGAA GACGCAGAAA	14520
TATTTAAAAC TCTTTAAAAT CTAATTTTA AAGAAGTCTT ACTCTGAGGG CCTATGCTG	14580
TAAAATAATG GGCTCTTTTT TGATGCCCAA AAGTGAGGTT TATATGAAAC AAGAATCAAC	14640
TGTTGATTTG TTAC	14654

(2) INFORMATION FOR SEQ ID NO: 107:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 6405 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 107:

AGAAAAATCT GCTTTACAGA AAATAAAAAT AATAGGAGAA AATCTATGTC AGATTTGAAA	60
AAATACGAAG GTGTCATTCC AGCCTTCTAC GCATGTTATG ATGATCAAGG AGAAGTAAGC	120
CCAGAACGTA CGCGTGCCTT GGTCAATAC TTCATGATA AAGGTGTTCA AGGTCTTTAT	180
GTCAATGGTT CTTCTGGTGA ATGTATCTAC CAAAGCGTTG AAGATCGCAA GTTGATTTTG	240
GAAGAAGTCA TGGCGGTAGC AAAGGTAAAT TGACCATTAT TGCCCATGTT GCTTGCAATA	300
ATACTAAAGA TAGTATGGAA CTTGCTCGCC ATGCTGAAAG CTTGGGAGTA GATGCTATTG	360
CAACGATTCC ACCAATTTAT TTCCGCTTGC CAGAATACTC AGTTGCCAAA TACTGGAACG	420
ATATCAGTTC TGCAGCTCCA AACACAGACT ACGTGATTTA CAACATTCCT CAATTGGCAG	480
GGGTGCTTT GACTCCAAGC CTTTACACAG AAATGTTGAA AAATCCTCGT GTTATCGGTG	540
TGAAGAACTC TTCTATGCCA GTTCAAGATA TCCAAACCTT TGTCAGCCTT GGTGGAGAAG	600
ACCATATCGT CTTTAATGGT CCTGATGAGC AGTTCCTAGG AGGACGCCTC ATGGGGGCTA	660
GGGTGGTAT CGGTGGTACT TATGGTGCTA TGCCAGAACT CTTCTTGAAA CTAATCAGT	720
TGATGCGGA TAAGGACCTA GAAACAGCGC GTGAATTGCA GTATGCTATC AACGCAATCA	780
TTGGTAAACT CACTTCTGCT CATGGAAATA TGTACGGTGT CATCAAAGAA GTCTTGAAAA	840
TCAATGAAGG CTTGAATATT GGATCTGTTC GTTCACCATT GACACCAGTG ACTGAAGAAG	900
ATCGTCCAGT TGTAGAAGCG GCTGCTGCCT TGATTCTGTA AACCAAGGAG CGTTCCTCT	960
AATCTAAAAG GAGGTATTTA TGACATATTA CGTTGCAATT GATATCGGTG GAACCAACAT	1020
CAAGTATGGT TTGGTTGATC AAGAGGGGCA ACTTCTTGAA TCGCATGAAA TGCCAACTGA	1080
GGCGCATAAG GGTGGACCTC ATATCTTACA AAAGACCAA GATATCGTAG CTAGTTATTT	1140
AGAAAAAGGC CCAGTAGCAG GTGTTGCCAT ATCTTCTGCT GGGATGGTGG ATCCGGATAA	1200

GGGTGAGATT	TTCTATGCTG	GGCCGCAAAT	CCCTAACTAC	GCAGGCACCC	AGTTCAAAAA	1260
GGAAATCGAA	GAAAGCTTTA	CTATTCCTTG	TGAGATTGAA	AATGATGTCA	ACTGTGCAGG	1320
TCTTGCTGAG	GCAGTATCTG	G TTCAGGCAA	GGGAGCAAGT	GTGACACTTT	GCTTGACCAT	1380
TGGAACCGGT	ATCGGTGGTT	GCTTGATTAT	GGATAGGAAA	GTCTTCCATG	GTTTTAGCAA	1440
TTCAGCCTGT	GAAGTCGGGT	ATATGCATAT	GCAGGATGGA	GCTTTTCAAG	ACTTGGCTTC	1500
TACAACAGCT	TTAGTGAAAT	ATGTAGCTGA	AGCCCATGGA	GAAGATGTTG	ATCAGTGGAA	1560
TGGCCGTAGA	ATTTTCAAAG	AAGCCACTGA	AGGAAACAAA	ATCTGCATGG	AAGGTATTGA	1620
CCGTATGGTT	GACTATCTAG	GAAAAGGTCT	GGCAAATATT	TGCTACGTTG	CCAATCCAGA	1680
AGTGGTTATT	CTTGGTGGTG	GTATCATGGG	GCAAGAGGCT	ATCCTCAAAC	CTAAGATCCG	1740
TACAGCCTTG	AAAGAGGCTT	TGGTACCAAG	TTTAGCAGAA	AAAACACGAT	TAGAATTTGC	1800
CCATCACCAA	AATACAGCAG	GGATGTTGGG	TGCATATTAT	CATTTTAAGA	CAAAACAATC	1860
CTAGTTTGGC	TCAGCCAAAC	TAGGATTTTC	TTACACGTTT	TTGTCTACGA	TAGCCGTTGA	1920
GTTTTTTTATT	TTCCCAGTAG	CTATTAAAGA	TTTTTTCCTT	GCTTTCGCGA	TTGATTTCCA	1980
AAAAGTAGGC	ATAAATCAAA	TCGATAAAGA	AGAGCATAGG	AAGTTGAGCG	GATATTCGTT	2040
GGATATAGGA	GGGTGGCTG	TGGGTGGCTA	CAAGAACAGT	CTCTGTATAG	GTCTGGCTAT	2100
CTTTATTGGG	AACACTTGTA	AAGAGTACAG	TCTTTGCCCC	CATCTCCTTA	GCATCTAATA	2160
GACTATCTAA	AATAGAAGGA	GTTGAGCCTG	AAAGTGAGAA	GCCCAGTACT	AGACAATTTT	2220
CATCCATGAT	GCTGGTTGTC	CAGGCAAAGC	CGTCTGGTTC	TGTCAAAGCT	TCGCAGACCA	2280
CACCTAGTCG	CATAAAACGT	AATTTTCAAT	CACGGGCGAC	GAGGCCAGAA	CTCCCTGTTC	2340
CAAAGAAGTA	GATACGCTCA	GCATCTTCGA	TTAGCTGGGC	AATTCGTTCT	AGTTGGATTT	2400
CGTCAATCAA	GTCTTGTGTT	TGTTCCCTCA	TATGCTATA	ACTTCTGAGG	ACTCGTTTGG	2460
TCAGTGGACT	GTGCTTGGAG	ACTTGGTTGG	CTTGATTTTC	TGCCTGATGT	TGGTATTGGA	2520
AAATAAATTC	TCGGTAGCCA	GTAAGCCAC	ACTTTTTTAGC	AAAGCGGGTC	AAAGCAGCTT	2580
GAGAAATATG	TAATTTTTGG	GTGACTTGT	GAGAAGATAA	ATCATCTGTA	ATCGTTTCAG	2640
CTTGCAAAAA	ATAGCGAGCG	ATTTCTTGT	CTAGGTCTGT	CATTTCTTCA	AAATGTGAAT	2700
CAATGATAGT	TGCGATATCT	GTTTGTCCA	TAGGGAAAGC	TCCTTTACAT	GAGTCATACT	2760
GGAAGACTAG	ATCAGAGAAT	AGTCACACTT	CATTATAACA	CATAATATAA	GGATAGATAA	2820
ATAAAAACGC	ATCTCTGTTT	TAAAAACGAA	AAAATCGAAA	AAGCTTCTCT	CTTTTCCATA	2880
ATTTTCTACT	CAAATTGTGG	TACAATTAAG	AGTAAGATTT	TAAGTTAGAA	ATGAGACTGA	2940

TTTGTATGAG	AAAATTTAAC	AGCCATTCTGA	TTCCGATTTCG	GCTTAATTTA	TTGTTTTCAA	3000
TCGTCATTTT	ACTCTTTATG	ACCATTATTG	GTCGTTTGTT	GTATATGCAG	GTTTTGAACA	3060
AGGATTTTTA	CGAAAAAAG	CTAGCTTCAG	CTAGTCAGAC	CAAGATTACA	AGCAGTTCAG	3120
CCCGTGGGGA	AATTTATGAT	GCTAGTGGAA	AACCTTTGGT	AGAAAATACG	TTAAAGCAGG	3180
TTGTTTCCTT	TACGCGTAGC	AATAAAATGA	CGGCTACAGA	CTTAAAAGAA	ACAGCTAAAA	3240
AGTTACTGAC	TTATGTGAGC	ATCAGTTCTC	CAAATTTGAC	AGAACGCCAG	CTGGCGGATT	3300
ACTATTTGGC	TGATCCTGAA	ATCTATAAAA	AAATAGTGGG	AGCTCTCCCA	AGTGAGAAAC	3360
GCTTGGATTG	AGATGGCAAT	CGTCTATCCG	AATCAGAACT	GTATAACAAT	GCGGTCGATA	3420
GTGTACAAAC	GAGTCAACTA	AACTATACAG	AGGATGAAAA	GAAAGAAATC	TATCTTTTTA	3480
GTCAGTTAAA	TGCTGTTGGA	AACTTTGCGA	CAGGAACCAT	TGCGACAGAT	CCTCTAAATG	3540
ATTCTCAGGT	GGCTGTTATT	GCCTCTATTT	CAAAGGAGAT	GCCTGGCATT	AGTATTTCTA	3600
CTTCTTGGGA	TAGAAAGGTT	TTGGAAACTT	CCCTTTCTTC	TATAGTTGGG	AGTGTATCCA	3660
GTGAAAAGC	TGGTCTCCCA	GCGGAAGAAG	CAGAAGCCTA	TCTTAAAAAA	GGCTATTTCTC	3720
TAAATGACCG	TGTAGGAACC	TCCTATTTGG	AAAAGCAATA	TGAAGAGACC	TTACAAGGAA	3780
AACGCTCGGT	AAAAGAAATC	CATCTGGATA	AATATGGCAA	TATGGAAAGC	GTGGATACAA	3840
TTGAGGAAGG	TAGTAAGGGA	AACAATATCA	AACTGACCAT	TGATTTGGCT	TTCCAAGATA	3900
GCGTGGATGC	TTTACTGAAA	AGTTATTTCA	ATTCTGAGCT	AGAAAATGGT	GGAGCCAAGT	3960
ATTCTGAAGG	TGTCTATGCA	GTCGCCCTTA	ACCCAAAAAC	AGGTGCGGTT	TTGTCTATGT	4020
CAGGGATTAA	ACATGACTTG	AAAACGGGAG	AGTTGACGCC	TGATTCCTTG	GGAACGGTAA	4080
CCAATGTCTT	TGTTCCAGGT	TCGGTTGTCA	AGGCGGCGAC	CATCAGCTCA	GGTTGGGAAA	4140
ATGGAGTCTT	GTCAGGAAAC	CAGACCTTGA	CAGACCAGTC	CATTGTCTTC	CAAGGTTGAG	4200
CTCCCAFCAA	TTCTTGGTAT	ACTCAGGCTT	ACGGTTCATT	CCCTATCACA	GCGGTCCAAG	4260
CTCTGGAGTA	TTCATCAAAAT	ACCTATATGG	TCCAACAGC	CTTAGGTCTT	ATGGGGCAAA	4320
CCTATCAACC	CAATATGTTT	GTCGGCACCA	GCAATCTAGA	GTCTGCTATG	GAGAAACTGC	4380
GTTCAACCTT	TGGCGAATAT	GGCTTGGGTA	CTGCGACAGG	AATTGACCTA	CCAGATGAAT	4440
CTACTGGATT	TGTTCCAAA	GAGTATAGCT	TTGCTAATTA	CATTACTAAT	GCCTTTGGGC	4500
AGTTTGATAA	CTATACGCCG	ATGCAGTTGG	CTCAGTATGT	AGCAACTATT	GCAAATAATG	4560
GTGTTTCGTG	GGCTCCTCGT	ATTGTTGAAG	GCATTTATGG	TAATAATGAT	AAGGGAGGAC	4620
TGGGTGACTT	GATTCAGCAA	CTGCAACCGA	CAGAGATGAA	TAAGGTCAAT	ATATCCGACT	4680
CCGATATGAG	CATCTGCAC	CAAGTTTTTT	ATCAGGTTGC	CCATGGTACT	AGTGGATTGA	4740

CAACTGGACG TGCCTTTTCA AATGGTGCCT TGGTATCCAT TAGCGGAAAA ACAGGTACAG 4800
 CCGAAAGCTA TGTGGCAGAT GGTCAGCAAG CAACCAATAC CAATGCGGTG GCCTATGCC 4860
 CATCTGATAA TCCCCAAATC GCTGTGCGAG TGGTCTTTCC TCATAATACC AATCTAACAA 4920
 ATGGTGTAGG ACCTTCCATT GCGCGTGACA TTATCAATCT GTATCAAAAA TACCATCCAA 4980
 TGAATTAGAA AGGAAATTAT GCTTTATCCA ACACCTATTG CCAAGTTGAT TGACAGTTAT 5040
 TCTAAGTTAC CAGGTATCGG GATTAAGACG GCTACGCGTC TGGCCTTTTA TACGATTGGG 5100
 ATGTCTGCTG ATGATGTCAA TGAATTTGCA AAAAAATCTCC TTCTGCTAA GAGAGAAATG 5160
 ACATATTGTT CTATTGTGG ACGTTTGACA GACGACGATC CTTGTTCTAT CTGTACTGAT 5220
 CCGACTCGTG ACCAGACAAC AATTTTAGTT CTGAGGATA GTAGAGATGT GGCAGCCATG 5280
 GAAAATATCC AAGAATACCA TGGACTCTAT CATGTCCTTC ATGGCCTCAT TTCTCCTATG 5340
 AATGGTATCA GTCCGACGA TATCAATCTC AAGAGCCTTA TGACTCGTCT TATGGATAGT 5400
 GAGGTTTCAG AAGTGATTGT GCGGACTAAT GCTACAGCGG ATGGTGAAGC GACTTCCATG 5460
 TATCTTTCAC GTTTGCTCAA GCCGGCTGGT ATCAAGGTTA CGCGTCTAGC ACGAGGTCTC 5520
 GCTGTGGGAG CGGACATTGA GTATGCGGAC GAAGTGACAC TCTTACGAGC CATTGAAAA 5580
 CGGACAGAGT TGTAAGTGTA GGCAAATTTA CGAACTCCAT TCATTTATAA AAAATCAAAG 5640
 AGGCTGAAAA TCGTTCCTAT CGGCCTCTTT TTGTATAGTG TGATGAGTAG GCTCAGGTT 5700
 AAGTTTAAA AAACCAAGCA AATATGATAT ACTAAAGAGC GAGTATTCTA GTAGAATTAG 5760
 GACAAATAAT ATGAAACAAA CGATTATCT TTTATATGGT GGACGGAGTG CGGAACGCGA 5820
 AGTCTCTGTC CTTTCAGCTG AGAGTGTCAT GCGTGCGGTC GATTACGACC GTTTCACAGT 5880
 CAAGACTTTC TTTATCAGTC AGTCAGGTGA CTTTATCAAA ACACAGGAAT TTAGTCATGC 5940
 TCCGGGGCAA GAAGACCGTC TCATGACCAA TGAAACCATT GATTGGGATA AGAAAGTTGC 6000
 ACCAAGTGCT ATCTACGAAG AAGGTGCAGT GGTCTTTCCA GTCCTTCACG GGCCAATGGG 6060
 AGAAGATGGC TCTGTCAAG GATTCTTGA AGTTTTGAAA ATGCCTTACG TTGGTTGCAA 6120
 CATTTGTCA TCAAGTCTTG CCATGGATAA AATCAGACT AAGCGTGTT TGGAACTGTC 6180
 TGGTATTGCC CAAGTTCCTT ATGTGGCTAT CGTTGAAGGC GATGATGTGA CTGCTAAAA 6240
 CGCTGAAGTG GAAGAAAAAT TGGCTTATCC AGTCTTCACT AAGCCGTCAA ACATGGGGTC 6300
 TAGTGTCGGT ATTTCTAAGT CTGAAAACCA AGAAGAACTC CGTCAAGCCT TAAAACCTGC 6360
 CTTCCGATAT GACAGCCGTG TCTTGGTTGA GCAAGGAGTG AATGC 6405

(2) INFORMATION FOR SEQ ID NO: 108:

794

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 11309 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 108:

CGAGCTCGGG TACCGGGATT TTAAGGAGTT TGATATGTAT AACCTATAT TAACCATTTT	60
ATTAGTATTA TCTGTTGTGA TTGTGATTGC AATTTTCATG CAACCAACCA AAAACCAATC	120
CAGCAATGTA TTTGATGCCA GTTCAGGTGA TTTGTTTGAA CGCAGTAAAG CTCGCGGTTT	180
TGAAGCTGTA ATGCAGCGTT TGACAGGGAT TTTAGTCTTT TTCTGGCTAG CCATTGCCTT	240
AGCATTGACG GTATTATCAA GTAGATAAGA AAATAATGGG CAGGACTAGG TCTTTCCTC	300
TTTTTATTTT TAAAGGATGT TTGAGAAGGT TTTACAGTAA AAGAAAATTA AAAAATCTAG	360
AAAGAAAATA TGAAAGATAG AATAAAAGAA TATTTACAAG ACAAGGGAAA GGTGACTGTT	420
AATGATTTGG CTCAGGCTTT GGGAAAAGAC AGTTCCAAGG ATTTTCGTGA GTTGATTAAA	480
ACCTTGTCCT TAATGGAAAAG AAAGCACCAA ATTCGTTTGG AAGAAGATGG TAGTCTGACA	540
TTAGAAATTA AGAAAAACA TGAGATTACC CTCAAGGGGA TTTTTCATGC CCATAAAAAAT	600
GGCTTTGGCT TTGTTAGTCT GGAAGGCGAG GAGGACGACC TTTTGTAGG GAAAAATGAT	660
GTCAACTATG CTATTGATGG TGATACCGTC GAGGTAGTGA TTAAGAAAGT CGCTGACCGC	720
AATAAGGGAA CAGCAGCAGA AGCCAAAATT ATTGATATCC TAGAACACAG TTTGACAACA	780
GTTGTCGGGC AAATCGTTCT GGATCAGGAA AAACCTAAGT ATGCTGGCTA TATTCGTTCA	840
AAAAATCAGA AAATCAGTCA ACCGATTTAT GTTAAGAAAC CAGCCCTAAA ATTAGAAGGA	900
ACAGAAGTTC TCAAAGTCTT TATCGATAAA TACCCAAGCA AGAAACATGA TTTCTTTGTC	960
GCGAGTGTTC TCGATGATG GGGACACTCA ACGGATGTCG GAATTGATGT TCTTGAGGTC	1020
TTGGAATCAA TGGACATTGT ATCCGAGTTT CCAGAAGCTG TTGTTAAGGA AGCAGAAAGT	1080
GTGCCTGATG CTCCGTCTCA AAAGGATATG GAAGGTCGTC TGGATCTAAG AGATGAAATT	1140
ACCTTTACCA TTGACGGTGC GGATGCCAAG GACTTGGACG ATGCAGTGCA TATCAAGGCT	1200
CTGAAAAATG GCAATCTGGA GTTTGGGGTT CACATCGCAG ATGTTTCTTA TTATGTGACC	1260
GAGGGTCTG CCCTTGACAA GGAAGCCCTT AACCGTGCGA CTTCTGTTTA CGTGACAGAC	1320
CGAGTGGTGC CAATGCTTCC AGAACGACTA TCAAATGGCA TCTGCTCTCT CAATCCCCAA	1380
GTTGACCGCC TGACCCAGTC TGCTATTATG GAGATTGATA AACATGGTCG TGTGGTCAAC	1440
TATACCATTA CACAAACAGT TATCAAGACC AGTTTTCGTA TGACCTATAG CGATGTCAAT	1500

795

GATATCCTAG	CTGGCGATGA	AGAAAAGAGA	AAAGAATATC	ATAAAATTGT	ATCAAGTATC	1560
GAACTCATGG	CCAAGCTTCA	TGAAACTTTA	GAAAACATGC	GTGTGAAACG	TGGAGCTCTC	1620
AATTTTGATA	CCAATGAAGC	GAAGATTTTA	GTGGATAAAC	AAGGTAAGCC	TGTTGATATC	1680
GTTCTTCGGC	AGCGTGGTAT	TGCCGAGCGG	ATGATTGAGT	CTTTTATGTT	GATGGCTAAT	1740
GAAACAGTTG	CCGAACATTT	CAGCAAGTTG	GATTTGCCTT	TTATCTATCG	AATTCACGAG	1800
GAGCCTAAGG	CTGAAAAGGT	TCAGAAGTTT	ATTGATTATG	CTTCGAGTTT	TGGCTTGC GC	1860
ATTTATGGAA	CTGCCAGTGA	GATTAGTCAG	GAGGCAC TTC	AAGACATCAT	GCGTGCTGTT	1920
GAGGGAGAAC	CTTATGCAGA	TGTATTGTCC	ATGATGCTTC	TTCGCTCTAT	GCAGCAGGCT	1980
CGTATTCGG	AGCACAATCA	CGGCCACTAT	GACTAGCTG	CTGACTATTA	TACTCACTTT	2040
ACCAGTCCAA	TTCGTCGTTA	TCCAGACCTT	CTTGTTCACC	GTATGATTCG	GGATTACGGC	2100
CGTCTAAGG	AAATAGCAGA	GCATTTTGAA	CAAGTGATTC	CAGAGATTGC	GACCCAGTCT	2160
TCCAACCGTG	AACGTCGTGC	CATAGAAGCT	GAGCGTGAAG	TCGAAGCCAT	GAAAAGGCT	2220
GAGTATATGG	AAGAATACGT	GGTGAAGAG	TATGATGCAG	TTGTATCAAG	TATTGTCAAA	2280
TTCGGTCTCT	TTGTCGAATT	GCCAAACACA	GTTGAAGGCT	TGATTCACAT	CACTAATCTG	2340
CCTGAATTTT	ATCATTTCAA	TGAGCGTGAT	TTGACTCTTC	GTGGAGAAAA	ATCAGGTATC	2400
ACTTTCGGAG	TGGGTCAGCA	GATCCGTATC	CGTGTGAAA	GAGCGGATAA	AATGACTGGA	2460
GAGATTGATT	TTTCATTCGT	ACCTAGTGAG	TTTGATGTGA	TTGAAAAAGG	CTTGAAACAG	2520
TCTAGTCGTA	GTGGCAGAGG	GCGTGATTCA	AATCGTCGTT	CGGATAAGAA	GGAAGACAAG	2580
AGAAAATCAG	GACGCTCAA	TGATAAGCGT	AAGCATTCAC	AAAAAGACAA	GAAGAAAAA	2640
GGAAAGAAAC	CTTTTACAA	GGAAGTAGCT	AAGAAAGGAG	CCAAGCATGG	CAAAGGGCGA	2700
GGGAAAGGTC	GTCGCACAAA	ATAAAAAGGC	ACGCCACGAC	TATACAATCG	TAGATACGCT	2760
AGAGGCAGGG	ATGGTCTTGA	CTGGAACTGA	AATCAAGAGT	GTACGAGCTG	CTCGAATTAA	2820
TCTCAAGGAT	GGCTTTGCTC	AAGTGAAAA	TGGAGAAGTT	TGGCTGAGCA	ATGTTTCATAT	2880
CGCGCCTTAC	GAAGAGGGCA	ATATCTGGAA	CCAGGAACCA	GAACGTCGTC	GTAAACTCCT	2940
GCTCCATAAA	AAGCAAATTC	AAAAATTGGA	ACAAGAGATC	AAAGGGACAG	GAATGACCTT	3000
AGTTCCCTT	AAGGTCTATA	TAAAAGATGG	CTACGCTAAG	CTTCTTTTAG	GACTTGCCAA	3060
AGGAAGCAT	GACTATGACA	AACGGGAGTC	TATCAAACGT	CGTGAGCAA	ATCGAGATAT	3120
CGCGCGTGTG	ATGAAAGCTG	TTAATCAGCG	ATAAAAAGAG	GAATTGAAAA	TGGAAAAATT	3180
AGTTGCCTAT	AAACGCATGC	CTTTGTGGAA	TAAACAAACA	ATGCCTGAAG	CTGTTACGCA	3240

AAAGCACAAT	ACAAAAGTTG	GGACTTGGGG	GAAAATTACT	GTCTTGAAGG	GAGCTCTCAA	3300
GTTTATTGAA	TTGACAGAAG	AAGGGGAAGT	TCTAGCTGAA	CACCTCTTTG	AAGCAGGGGC	3360
AGACAATCCA	ATGGCCCAAC	CTCAAGCCTG	GCACCGAGTG	GAAGCTGCCA	CAGATGATGT	3420
GGAATGGTAC	TTGGAATTTT	ATTGTAAACC	TGAGGATTAT	TTTGCTAAAA	AATACAATAC	3480
CAATCCTGTT	CATTCAGAGG	TCCTAGAGGC	CATGCAGACA	GTGAAACAAG	GGAAAGCTTT	3540
GGATTTGGGT	TGTGGTCAGG	GGCGTAATTC	TCTTTTCTA	GCCCAGCAAG	ATTTTGATGT	3600
GACGGCTGTA	GATCAAAATG	GACTIONCTCT	TGAAATCTTG	CAAAGCATTG	TGGAGCAGGA	3660
AGATTTGGAC	ATGCCTGTTG	GCCTTTACGA	TATCAATTCA	GCTAGCATTG	AACAAGAATA	3720
TGATTTTATC	GTTTCAACAG	TTGTTCTCAT	GTTTCTACAA	GCGGACCGCA	TTCCAGCTAT	3780
TATTCAAAAT	ATGCAGGAGA	AAACCAGTGT	TGTTGGTTAC	AACCTTATCG	TTTGTGCCAT	3840
GGACACGGAG	GATTATCCTT	GCTCGGTTAA	CTTCCCATTG	ACCTTTAAAG	AAGGAGAACT	3900
GGCAGACTAT	TACAAGGATT	GGGAATTGGT	TAAGTACAAT	GAAAATCCAG	GCCATTTGCA	3960
CCGTCGCGAT	GAGAATGGCA	ATCGTATTCA	ACTACGCTTT	GCGACCTTAC	TAGCTAAGAA	4020
AATCAAGTAA	ACACACATGA	AGATTAGGAA	TTTTCCTGAT	CTTTTCTCTT	TTTACGAAT	4080
GATATAGAAA	AGGAGGGAAT	TCATGTTTGT	TGCAGAGAT	GCTAGGGGAG	AATTGGTAAA	4140
TGTGTTAGAG	GATAAACTTG	AGAAGCAAGC	ATACACCTGC	CCAGCTTGTG	GAGGCCAGCT	4200
CCATTTGCGT	CAAGGACCAA	GTGTACGGAC	GCATTTTGCC	CATAAATCCT	TAAAAGACTG	4260
TGATTTTTTC	TTTGAAAATG	AAAGTCCAGA	ACACCTGGCC	AATAAGGAAT	CCCTCTATCA	4320
CTGGTTGAAA	AAAGAGACAA	AGGTTCAATT	AGAGTACCCG	CTTTCAGAAC	TTAAACAGAT	4380
TGCGGATGTA	TTTGTAATG	GCAATCTAGC	TCTAGAAGTT	CAGTGTAGTC	CCTTGCCCTCA	4440
GAAAGTCCTT	AAAGAGCGAA	GTGAGGGCTA	TCGTAGTCAG	GGTTACCAAG	TACTGTGGTT	4500
GCTGGGTCAA	AAACTGTGGC	TCAAGGAGCG	TTTGACTCGT	CTACAGCAAG	GTTTTCTTTA	4560
TTTCAGTCAA	AACATGGGCT	TTTATGTTTG	GGAATTAGAC	AAGGAAAAAC	AAGTTTTAAG	4620
ACTCAAATAC	CTGATTTACC	AGGATCTCCG	CGGTAAACTC	CATTATCAAA	TCAAGGAATT	4680
TTCTATGGT	CAAGGTAGTT	TATTGGAAAT	ATTGCGTCTT	CCCTATAAGA	GACAAAAAAT	4740
ATCTCATTTT	ACAGTTTCTG	AGGACAAGGA	CATCTGTGCG	TATATCCGGC	AACAACTTTA	4800
TTATCAAAAT	CTCTTTTGGG	TGAAAGAACA	AGCAGAAGCC	TATCAAAAAG	GAGAAAATAT	4860
CCTGACTTAT	GGACTGAAAG	AATGGTATCC	ACAAATTCGA	CCAATAGTGG	GCAAATTTTT	4920
CCAGATTGAA	CAAGACTTGA	CTAGCTATTA	TCAGCACTTT	TATACCTATT	ACCAAAAAAA	4980
TCCTCAAAAT	GATTGGCAA	AGCTTTATCC	ACCAGCCTTT	TATCAGCAAT	ATTTCTTGAA	5040

797

AAATATGGTA GAATAGAAAG GATGGAGGAA TCTAATGGTA TTACAAAGAA ATGAAATAAA	5100
TGAAAAAGAT ACATGGGATC TATCAACGAT CTACCCAAC T GACCAGGCTT GGGAAGAAGC	5160
CTTAAAAGAT TTAACAGAAC AATGAGGAC AGTAGCCCAG TATGAAGGCC ATCTCTTGGA	5220
TAGTGCGGAT AACCTACTAG AAATCACTGA ATTTTCTCTT GAAATGGAAC GCCAGATAGA	5280
GAAGCTTTAC GCTTATGCTC ATATGAAGAA TGACCAGGAT ACACGTGAAG CTAAGTATCA	5340
AGAGTACTAT GCCAAGGCCA TGACACTCTA CAGCCAGTTA GACCAAGCCT TTTCATTCTA	5400
TGAGCCTGAA TTATGGAGA TTAGCGAAAA GCAGTATGCT GACTTTTTAG AAGTCAACC	5460
AAAGCTGCAG GTTATCAAC ACTATTTTGA CAAGCTTTTG CAAGGCAAGG ATCACGTTCT	5520
TTCAACAACGT GAAGAAGAAT TATGGCTGG AGCTGGAGAA ATCTTTGGTT CAGCAAGTGA	5580
AACCTTCGCT ATCTTGACA ATGCGGATAT TGTGTTCCCT TATGTCCTAG ACGATGATGG	5640
TAAAGAAGTT CAGCTATCTC ATGGGACTTA CACACGTTTG ATGGAGTCTA AAAACGTGA	5700
GGFTCGCCGT GGTGCCTATC AAGCTCTTFA TCGGACTTAC GAACAATTCC AACACACCTA	5760
TGCCAAAACC TTGCAAACCA ATGTTAAGGT GCAAAATTAC CGTGCTAAAG TTCGTAACCTA	5820
CAAGAGTGCT CGTCATGCAG CCCTCGCAGC GAATTTTGTT CCAGAAAGTG TTTATGACAA	5880
TTTGGTAGCA GCAGTTCGCA AGCATTTGCC ACTCTTACAT CGTATCTTG AGCTTCGTTT	5940
AAAAATCTTG GGGATTTTCA ATCTCAAGAT GTACGATGTC TACACACCGC TTTCATCTGT	6000
TGAATACAGT TTTACCTACC AAGAAGCCTT GAAAAAGCA GAAGATGCTT TGGCAGTCTT	6060
GGGTGAGGAT TACTTGAGCC GTGTTAAACG TGCCTTCAGC GAGCGTTGGA TTGATGTTTA	6120
CGAAAATCAA GGCAAGCCTT CAGGTGCCTA CTCTGGTGGT TCTTATGATA CCAATGCCTT	6180
TATGCTTCTC AACTGGCAAG ACAATCTGGA CAATCTCTT ACTCTTGTTT ATGAAACAGG	6240
TCACAGTATG CATTCAAGCT ATACTCGTGA AACTCAGCCT TATGTTTACG GGGATTACTC	6300
TATCTTTTTG GCTGAGATTG CCTCAACTAC CAATGAAAAT ATCTTGACGG AGAAATTATT	6360
GGAAGAAGTG GAAGACGACG CAACACGCTT TGCTATTCTC AATAACTTCC TAGATGGTTT	6420
CCGTGGAACA GTTTTCCGCC AACTCAATT TGCTGAGTTT GAACACGCCA TTCACCAAGC	6480
AGATCAAAAT GGGGAGGTCT TGACAAGCGA TTTCCCTAAAT AACTCTACG CAGACTTGAA	6540
CCAAGAGTAT TATGGTTTGA GTAAGGAAGA CAATCCTGAA ATCCAATACG AGTGGGCTCG	6600
CATTCCACAC TTCTACTATA ACTACTATGT ATATCAATAT TCAACTGGCT TTGCGGCCGC	6660
CTCAGCCTTG GCTGAAAAA TTGTCCATGG TAGTCAAGAA GACCGTGACC GCTATATCGA	6720
CTACCTCAAG GCAGGTAAGT CGGACTATCC ACTTAATGTC ATGAGAAAAG CTGGTGTGTA	6780

798

TATGGAGAAG GAAGACTACC TCAACGATGC CTTTGCAGTC TTTGAACGCC GTTTAAATGA 6840
 GTTTGAAGCC CTTGTTGAAA AATTAGGATTT GGCATAAAAT GGTGAATCG TATAGTAAGA 6900
 ATGCTAACCA TAACATGCGT CGTCCTGTCG TCAAAGAAGA AATTGTAGAC TTGATGCGTC 6960
 AGCGTCAAAA GCAGGTCACA GGTTCCTTGA AAGAATTGGA AGACTTTGCC CGCAAGGAAA 7020
 ATATTCCTAT TATFCCCAT GAAACGGTTG CTTATTTCCG TTTTCTTATG GAAACCATGC 7080
 AGCCTAAAAA TATFCTGGAA ATTGGGACGG CTATCGGTTT TTCAGCTCTC TTGATGGCTG 7140
 AACATGCGCC AAATGCTAAG ATTACAATA TTGATCGTAA TCCAGAAATG ATTGGTTTGT 7200
 CCAAGGAAAA TTTTGCCAG TTTGACAGTC GCAAGCAAAT CACTCTCCTA GAGGGAGATG 7260
 CGGTGGATGT CTTATCTACA CTGACAGAGT CTTATGATTT CGTCTTTATG GATTCTGCCA 7320
 AGTCTAAATA CATCGTCTTT CTGCCAGAAA TCCTCAAACA TTTGGAAGTT GGTGGTGTGG 7380
 TTGTCTTGA TGATATTTT CAAGGTGGTG ATGTTGCCAA GGATATTATG GAAGTCCGTC 7440
 GTGGTCAGCG AACCATTTAT CGAGGCCTC AAAAATTATT TGATGCAACC TTAGACAATC 7500
 CAGAACTCAC CGCAACATTA GTGCCTTTAG GAGATGGTAT TCTCATGCTT CGTAAAAATG 7560
 TAGCAGATGT TCAACTGTCT GAAAGCGAAT GATTTTCAGA AAAATTTAAG AAAAAATAGT 7620
 AAAATAGATA GAGTAACACT TATCTCAAAG GAGTAGACAT GAAGAAAAA TTATTTGGCAG 7680
 GTGCCATCAC ACTATTATCA GTAGCAACTT TAGCAGCTTG TTCGAAAGGG TCAGAAGGTG 7740
 CAGACCTTAT CAGCATGAAA GGGGATGTCA TTACAGAACA TCAATTTTAT GAGCAAGTGA 7800
 AAAGCAACCC TTCAGCCCAA CAAGTCTTGT TAAATATGAC CATCCAAAAA GTTTTGTAAA 7860
 AACAATATGG CTCAGAGCTT GATGATAAAG AGGTTGATGA TACTATTGCC GAAGAAAAA 7920
 AACAATATGG CGAAAACACT CAACGTGTCT TGTACAAGC AGGTATGACT CTTGAAACAC 7980
 GTAAAGCTCA AATTCGTACA AGTAAATTAG TTGAGTTGGC AGTTAAGAAG GTAGCAGAAG 8040
 CTGAATTGAC AGATGAAGCC TATAAGAAA CTTTGTGATGA GTACACTCCA GATGTAACGG 8100
 CTCAAATCAT CCGTCTTAAT AATGAAGATA AGGCCAAAAGA AGTTCTCGAA AAAGCCAAGG 8160
 CAGAAGGTGC TGATTTTGCT CAATTAGCCA AAGATAATTC AACTGATGAA AAAACAAAAG 8220
 AAAATGGTGG AGAAATTACC TTTGATTCTG CTTCAACAGA AGTACCTGAG CAAGTCAAAA 8280
 AAGCCGCTTT CGCTTTAGAT GTGGATGGTG TTTCTGATGT GATTACAGCA ACTGGCACAC 8340
 AAGCCTACAG TAGCCAATAT TACATTGTAA AACTACTAA GAAAACAGAA AAATCATCTA 8400
 ATATTGATGA CTACAAAGAA AAATTAATA CTGTTATCTT GACTCAAAA CAAAATGATT 8460
 CAACATTTGT TCAAAGCATT ATCGGAAAAG AATTGCAAGC AGCCAATATC AAGGTTAAGG 8520
 ACCAAGCCTT CAAAATATC TTTACCCAAT ATATCGGTGG TGGAGATTCA AGCTCAAGCA 8580

GTAGTACATC	AAACGAATAG	TCCAAATCAA	TGAGTCAGGG	AAAAAECTCG	ACTTCAGGAA	8640
AAAATGAAGC	AAACATTCCC	ACAATAAAAC	GCATAGTACA	AGGTTTGTAC	TGCCCCCAA	8700
AAAGTTAGAC	AATTAATTTA	TCCGAAGGAT	TTAGTTCTGT	ATTGCACAGA	GCTAAGTCCT	8760
TTTAGTTTTA	TCTTAATTCCT	CTTATTGTTG	TAATAATCAA	TATAGTCTAT	AATGGCTCGT	8820
TCCAATTGAT	TAAGTGATTT	AAATGTTTTTC	TCATAGCCAT	AAAACATTTT	GGATTTTAAA	8880
ATGCCAAAGA	AAGATTCCAT	CCTACCGTTG	TCTTGGCTGT	TGCCCTTACG	TGACATGGAT	8940
GCTTGAATTC	CCTTACTCTC	TAGGAAGCGA	TGATAAGAAT	CGTGTGATA	TTGCCAGCCT	9000
TGGTCACTAT	GGAGAATCGT	ATTCTCGTAG	TGCTTCTCTT	TGAATGCCTG	TTCCAACATT	9060
AACGATCAAT	CAATTTAATC	ATGTACCTAA	GATTAGAATT	GTTTATCCCA	AATTTATTTG	9120
AAAGCTTCTC	TAAGCTATAT	CCTTGTTTTC	TAAGTTCATA	GATCTGAACT	TTATCATCAT	9180
AAGTTAATTT	CATAATAAAA	ACACCCCAA	AGTTAGATTT	TTTCTGTCTA	ACTTTTGGGG	9240
TGTAGTTCAT	GTACACCTGA	TATGATGCGT	TTTATAATTT	TAAAGACTTT	TTGACCAGCC	9300
TCATTTTTTT	AACTTGATAC	TCAGTGA AAA	GCAAAGATTA	AACTAGGAAG	CTAGCTGTAG	9360
GCTGCTCAA	GAACAGCTTT	GAGGTTGTAG	ATAAACTTG	TGAGGTCACC	AACATATATA	9420
ATGTGAAGCT	GACGTGGTTT	GAATAGATTT	TAGAAGAGTA	TGAGTCTGGA	AGTTTTAATG	9480
GATAATGCAA	GATTCCATAG	AATGGGTAAG	CTAGAGTTCT	TATGTGAAGA	GTTTGGGCAT	9540
AAACTTTTAC	CTTTTCCCTC	CTACTCATCT	TAGTATAGAA	AAGTGAATCT	GAAATAGTAC	9600
ATAACTGCTT	CTAAAACATT	CTTATAAATT	GATTTAAATT	CTCAAATCAT	ATTATTCAGT	9660
TCTTATTTCA	TTTTGTTCCTA	CAATCCTGTT	GAGAAGACAC	GTGTTTATAT	CAAAAAGGTA	9720
TTGGCAAGTT	GCAATACCTT	TTTACGAGGC	TCTGTTGTCT	TATTTTTGTT	TCAACTGACT	9780
ATATCTCCTA	TGGTCTTAGT	TCAGAAGGCT	AGGCTATAAT	TATGATTGAT	AAGAAGTATC	9840
ATTCCAAGTA	TTGGGAGTGA	ATGTTTCAA	ATCATGGGTT	TCTATAATGG	TCAGGCTGGC	9900
ATTTGCTAGA	CCGCCATCTT	TACGAAGAAG	TGGTTCTTTA	TAGCCTAGGA	GAGTACGAAG	9960
ACTGGCAGTA	AGATTGGCGC	CGTGTCCGAC	AATTAGAATA	CGTTCAGCTG	GACTATCTTT	10020
TAATGATTTG	ATAAATTGGA	TGGTCCGTTG	AGTTGTACTA	TAGAGGGATT	CGGCTCCGAA	10080
CATTCGAGTG	TCAAATTGAG	CAAGATTTGA	ACGAAAAGCC	TGGATTTGTT	GCGGGTAAAT	10140
AGCTTCCAAG	GTGCAATTT	TCAAACCTTC	TAACCTCCCA	AGTTGCCATT	CACGGAGATT	10200
AGGAACGATT	TCTAAAGAAC	AGGGGGTATA	GAGTTGACTT	TGGATAATCT	CAGCAGATTT	10260
GACCGCTCGA	GGTAAATCAC	TTGAATAAAT	CTGATCAAAA	GGAATTCCTT	TGAGATACTG	10320

800

ACCAAGTCGT TTTAGGGTTT CAATGGATTC AGGAAGAAGA GGAGAATCAC CACTAGCACC 10380
 TTGAAAACGA CCTTCTTGGT TCCAGAGGGT ACGACCGTGG CGGACAAAGT AGAGTTTCAT 10440
 TACTTGATGT CCTCCAAAAT ATCTACAAAG TCTGCCTTTA CAAAGCTAGC CAAGTCTTGT 10500
 GGC CGCAGCA TAATGCTGTG TCCGACTTCG CCTGCAGAGA CAATCATTTG ATCCAAATCT 10560
 AGAGCAATTT TATCGATAAA AATGGGATAA TTGTGTTTCT GACGAATTCC GACAGGATTA 10620
 TTGGCTCCAT GAATGTAACC AGTTGTTTTT TCTAAGTCCT TTTGTGGAAT CATGCTCACT 10680
 TTTTTATGTC CAGAAATTTT AGCTAGTTTC TTTTCAGACA AGTGCTGAGT GATAGGGACA 10740
 ATTCCGATAA TCGGTCCGGT CTTGTCTCCC AAAAGCGCCA AGGTTTTGAA AATCTGATCT 10800
 CGTTCATAAC CTFGAGGAAG CTCTCCTTCT AGGGCATTGA TTTGAATCCC CTGATGAGGG 10860
 ATAGCTGCTT TAGATAGGAT TTGTTCCACC AATGTTTTTT TGATTTTAAC TTTTTTTGCC 10920
 ATTATTTATA TTTATCCTCC AATTGACTCA TCCAAATACC AAGCCAGATT CCCAGCGCAA 10980
 AGAAGAAGGC GATGATGACA TAACCGACAA GTGAAAGTCC TGTGTATTGG ATACTTTCAG 11040
 CGTTTCCTGC ATTTGGAATT AAGATCAAAA GGGTACTTGA TAGGACGATA CCGATGATGA 11100
 AATGATAGAC GAACTGTTA CGGAGTTCTT CTAGTCTCC GTCCGTCAA GCGTAGGCCA 11160
 CTTCTCTTTT CTGCTTTA CCTTTGGACA TCTTGTAAG AGGTGGGAGG GCAATATAGA 11220
 CATGACCTGC CTCGACTAGC GGACGCATGT AACGGTAGAA AAATGTCAAG AGCAAGGTCT 11280
 GGATATGGGC ACCGTCGGTA TCCGCATCG 11309

(2) INFORMATION FOR SEQ ID NO: 109:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 5548 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 109:

CCATAGTCTA ACAAGTCTTT GTAAAGGTTT ATCCCTGATT CATGTAAAGA TTGTGTAAAG 60
 AATCAAAAAA AGCCACTTTT GAAAAATGGC TGCTCCTAAA AATAGCTTTA AAAATTATTA 120
 GTCCTGTGCG AAAGATFGGT TAGGAAGAAA AATCGTGAAG CAACTGCCTC TGCCAAGCTG 180
 ACTCGTCACC GTGACTTGGC CACCTAATAA TTGACTGAGT TCTTTGACAA TGGCAAGGCC 240
 AAGACCAGTG CCACCAGTTT GTCTGCTTCG ACCTTTATTA ACTCGGTAAA AACGTTCAAA 300
 AATACGATCC TGCTCTAATT GACTAATACC AATCCCTGTA TCTGATACAG AAATCTTAAT 360
 GCCTTCGTTT ACCTTTTGGG TCTTGACCTC AATTTTCCC CCTTGTTTCAG TGTAACGGAT 420

GGCATTGGAT AAAAGATTGA GTAAGATTTG GGAAAGTAAT TGA CTATCTG ATACGAGGGT	480
GACATCATCT GGCACCTGCA CCTTTAGCTG TAAATCCTTC TTCTTGAGCT GAGGTTGCAA	540
GCTTTGAGTC AAATCCTGTA CAAATCTGTC CAAAGAAAGG GTCGTCCATF GTATAGGCAT	600
TTGTTGAGCC TTAGATAAGG TAAGAAGATG CTCAACAATA TGCTCAAGAC GCAAACCTTC	660
TTTGTAATA ATGTCTAGAA AGTCATCCTT GAGCGCTTCT TCTTCAGCTG ACATCCCCTT	720
AATGGTTTCA GCAAAGCCCT TAATCGAAGT AACTGGTGTC CTCAATTCAT GGGAGGCATT	780
TGAGACAAAG GCTAAATTTA ACTTTTCATA AGTTCTAATC GTTGTTAAAT CATATAGCAA	840
GACGAGCACA GCTTCCACAG ATTGGGTGGG GCTAAAAACG GGAAGTGTGTC TCACTTCTAA	900
AATCAAGTCA CCCTCATGAA ACCCACTTAC TTCTTGTTTT AACCTTGTTT TTTGATCAAA	960
GGCTTGGTGA ACTAAATTC GAATATCCAT CCGTTTGAGG TCATCAAGTG AACTTATGTC	1020
GCCGTCCACA TCGGAAAAT AATGAGGCAG AGAGCGACTG GATAATAACA TCTGACCTTG	1080
AGCGGAAACT AAAAACGTCC CCATGGTTAG GTGCGACAGA AGAACCTCCA TTGTTTCGGC	1140
TAGATCCTTG TATTGCTGAT CCTGTTGGGA GACTTTGGTT TTTAGGCCAG ACACATACTG	1200
AGCCAAAGAC TTTAAGTCTT CTTGCCCTTT TTCTAAAAAG TATFCACTAC TGGTCAAGAG	1260
AGGTTGGTGC AAGGTCTCAA AAGCAACTTC CCATTTCCAA AGGCAAAGA GCCAGTAGCC	1320
ACCTAGTCCC AAAGAAAGGG CTAGAAGAAA GAGACCGATG CCTTACTGA TCCAAGTTAA	1380
TGCCATCCCT GCAATCAGAA TGAGGCTAAC ACTTAGATTG ACTAGCCAAA ATTGAAGGTA	1440
GCGTTTCATC TATAACTCCT TGAACCTATA ACCATAACCC CGAATGGTTC GAATAAATTG	1500
AGGGGCTTTA GGATTGTCTT CAATTTTTTC CCTCAACTTA CCAATATGAA CGTCCACCAA	1560
ACGTGTTTCC TGCCCAAAGT CATACCCCA GATACGTTCC AAAAGACGCT CTCTAGTCAG	1620
TGTCATGTTG GGATGTTTCA TAAGATAGAG CAAGAGTTCA AATTCCTTTG GGGTCAAAC	1680
CAGTAACTTA TTCGCCTTGT AGACTTCATG ACGCTCAGGG TATACTTTCA AGGTCCCAA	1740
TAGCCAAGAA TCGTCAGCGA TATTATCTGA ATCATCTCCT TCTTGTTCTC CTTTAGTTCTG	1800
CCTGAGGACA GCCTTGACAC GCGCCAGCAA TTCTCTAGGG CTAAAAGGCT TGTCAGGTA	1860
GTCATCAGCC CCTAATCCA AGGCCAAAAC CTTATCAAAT TCATCACTTT TCGCAGAAAC	1920
CATCATAATT GGAGTTTGA CGCCTTTGGC TCTCAGCCGC TTACAAACTT CCATGCCATC	1980
TAATTGTGGT AACATGATAT CAAGCAAGAT AAAATCAAAG GGTCTGTTT CTGCCAAAGC	2040
TAAGGCCTTC CGTCCATTTG TCACCAATTG AGTAGAAAAG CCTTCCTTAC TTAAATGGTA	2100
GTCAAGCAAT TTCAGAATGT GTTCTTCATC ATCCACTAAT AAGACTTGTT TTGTCATCTA	2160

802

TTATCTCCTA	TTGGTAACAT	TATAACACAA	TTATCAGAAA	TCCTAACATT	GCTAAATCAG	2220
ATTAAATTTG	CCTATCAAGA	CTAGTATCTG	GTCAAACGCT	CAATCATCTC	CTTGTGCTCT	2280
GGATAGGTCG	CCAGTAGATC	TACCCTTTCA	AATAATTCAA	AATCCTCAA	TTCAAAACCA	2340
GGAGCAACAA	GACAAGAAAC	CAGAGCATCA	TCCTTATCAA	CTGTTGATCC	CCAAATAGTG	2400
CCCTTAGGAA	CACAGTAGTG	AAGTTGTTGC	CCTTTGGATA	TGTCCAGGCC	TAAAGTGACT	2460
GCTTCGTAGT	GACCATCTGC	TGTAATCATG	TGAACAGTAA	GTGGGGATCC	TGCATGAAAA	2520
TACCAGATTT	CATCTGCTGT	CAATCGGTGA	AAATGTGAAG	GATTCGTTTC	TTCTAATAAG	2580
AAATAAATAC	TGGTATAAAG	CGCCCTTCCC	TTACCAGCAA	GGTTTATAGT	GTCTGAAGCT	2640
TTTTTTGTTT	GTCTAAAATA	GCCACCTTCA	ATATGGGGAG	CTAACTCTAG	AGTTCTTATC	2700
AAGTCTTCTT	TATCCGTCGG	AGCCAATGGG	TTGAAGTAAC	TCTTGTTCAA	AGTGGTTTTA	2760
CGATTTCAAG	AACTCCTCTC	AGTTCTGAGG	ACACGGTAAT	GATTGATGCG	ACGGAAGTAC	2820
AAATCAATCG	CCCTAAAAAA	AGAATTAGCG	AATGATCTCG	GTAAAAAAA	TGCCACGCTA	2880
TGAAGGCTCA	AGCGATTGTC	ACAAGTCAAG	GGAGAATTGT	TTCTTTGGAT	ATCGCTGTGA	2940
ACTATTGTCA	TGATATGAAG	TTGTTCAAAA	TGAGTCGCAG	AAATATCGGA	CAAGCTGGTA	3000
AAATCTTGGC	TGACAGTGGT	TATCAAGGGC	TCATGAAGAT	ATATCCTCAA	GCACAAACTC	3060
CACGTAAATC	CAGCAAACCTC	AAGCCACTAA	CAGTTGAAGA	TAAAGCCTAT	AACCATGCGC	3120
TATCCAAGGA	GAGAAGCAAG	GTTGAGAACA	TCTTTGCCAA	AGTAAAAACG	TTTAAAATGA	3180
TTTCAACAAC	CTATCGAAAT	CATCGTAAAC	ACTTCGGATT	ACGAATGAAT	TTGATTGCTG	3240
GCATTATCAA	TCATGAACTA	GGATTCTAGT	TTTGCAGGAA	GTCTATTATT	TGGTTAGGTG	3300
AATTAGTGAA	GCGTTTAGGC	AAGTGTCTCT	GGTTACGACG	TCATGGACTC	TAAATCGATT	3360
ATATTTAGGG	GTCATGACTA	GTGAAGCAGT	TAGCTAGTTC	GCATATAAGC	GGCTAGCGTC	3420
TAACAATTAG	GAACTTTAGT	TCCAATAACT	TTAAGATTAC	GACGTTTIFAG	GACATAAATC	3480
GATCATATTT	ATGTCCTAAA	ACTAGTGAAG	CGCCTAGCCA	AAGTCCGAAT	AGGATTTGGC	3540
GTTAGTFACT	TAGATGCTTT	TGCAATCAAG	TAACTTTGGC	GATTTACATC	TTCTCTGGCG	3600
CTTCTACTCC	AAGCAAGCGA	AGGGCTTCTT	TGAGAACGAC	TGCGGTTGCG	TAGCTGAGGG	3660
CTAGACGGCT	GTCGCGTTCT	GGGCTTTCAT	CCAAGATACG	TGTATGTGCA	TAGTATTTGT	3720
TAAAGGATTG	AGCCAGGCTA	ATTGCAAATT	TAGCAATGAT	AGAAGGTCA	AAGTTATCTG	3780
CCGCACGGTT	GATAATACGT	GGGAAGTCTT	GAATGAGTTT	AATGATTTCC	CAGCTTTTCAG	3840
TATCATTCAA	GCTATAGTTG	CCAGCTGTTT	CTGGTTTGAA	ATCGGCTTTG	CGTAAGATAG	3900
ATTGGATACG	AGCGTAGGCA	TATTGAACGT	AAGGTCCAGT	TTCACCTCG	AAGGATACCA	3960

TAGCCTCTAG GTCGAAGTCG TATCCATTTG TACGGTCGGT TTTGAGGTCA TAGAATTTAA 4020
 TGGCTCCAAT CCCAACAGCA TGTGCTACTT GGTCTTTGTT TTCTAGTTCA GGATTTTTAG 4080
 CCTCGATTTG GACCTTGGCA CGGCTAACAG CCTCTGCAAC AGTAGGCTCT AGCAAGATGA 4140
 CATTCCTTT ACGAGTAGAG AGTTTCTTCC CTTCTTTTGT AACCAAACCA AAAGGAACGT 4200
 GAGTAATGTC GTCACTCCAG TCGTAGCCCA TCTCTTGCAA GACAGCTTTG AGCTGTTTAA 4260
 AGTGGGCAGA TTGTTCTTGA CCAACGACAT AGATAGATTT AGCAAATTGG TATTCGTTTT 4320
 TACGGTAGAG GGCTGCAGCC AAGTCACGTG TGATATAGAG AGTTGCACCA TCAGACTTCT 4380
 TGATGAGGGC TGGATGTTCA ATTCCATATT TCTCAAGATT CACAACCTGG GCACCTTCTG 4440
 ATTCAAGAAG TAGTCCTTTT TCAGAAAGAA TGTCTACAAC TGCATCCATC TTATCATTTG 4500
 AGAAGGCTTC TCCGTTATAG CTGTCAAATT CAACCTTCAA TTCATTGTAA AGGCGGTAA 4560
 ATTCCACTAA ACTTTCATCG CGGAACCATT GCCAAAGAGC GAGAGCTTCC TCATCTCCAT 4620
 TTTCAAGTTT ACGGAACCAT TCGCGCGCTT CTTTATCCAA GCTAGGGTCA TTTTCAGCTT 4680
 CAGCGTTGAT GCGGACATAG AGTTTAAGGA GTTCATCGAT TGGATGAGCT TTTACAGCTT 4740
 CTTCGTCGCC CCATTTTTTG TAGGCAACAA TCAACATCCC AAATTGTTTA CCCAGTCTC 4800
 CCAAATGGTT GACCTTGACC GTTTGATAAC CGATTTTTTG GAAAATATGT GACAAGCTAT 4860
 CTCCGATAAC AGTTGAACGC AGGTGGCCAA TAGAAAATGG TTTAGCGATA TTCGGACTAG 4920
 ACATGTCGAT AACAAATTT TCTTGTTTAC CAATATTTTG GTCAGCATAG TGTCTTTTT 4980
 CAGTGGTAAC AGCTTGCAAT ACTTGAGCAG AAATGGCAGA TTTATCAAGG AAAAAGTTAA 5040
 CGTAAGGTCC TGTGCGACA ACTTTTTCAA AGGCTTGGCT GTTCATTTTT TCAGCCAGTT 5100
 CAGCCGCAAT CATTTGTGGT GCTTTACGTT CGACTTTTGC AAGAGAAAAA GCAGGAAAG 5160
 CAATGTCTCC CATTTCTGAG TTTTATAGGG TTTCCAGTAA CTTTAAAAATA GCCTCTTGGT 5220
 CCAGGCTATC AATGATGCTA GATAATTCGC TAGCAATCAA TTCTTTTGTA TTCATTAAGA 5280
 GTCCTTTTTT GGACTTTTCT ACTATTTTAT CACAATTTTA AAGAAAGAAG AAAAAATTTT 5340
 TGAAATCTCC TGTTTTTTTG GTATAATATG GTTATAAATA TAGTTATAAA TATGCACGCA 5400
 AGAGGATTTT ATGAGAAAAA GAGATCGTCA TCAGTTAATA AAAAAATGA TTAAGGAGGA 5460
 GAAATTAAGT ACACAAAAAG AAATTCAGA TCGGTTGGAG GCGCACAATG TTTGTGTGAC 5520
 GCAGACAACC TTGTCTCGTG ATTTGCGG 5548

(2) INFORMATION FOR SEQ ID NO: 110:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 3132 base pairs

804

(B) TYPE: nucleic acid
 (C) STRANDEDNESS: double
 (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 110:

TACCCGGTAG TCTTAGCAGA CACATCTAGC TCTGAAGATG CTTTAAACAT CTCTGATAAA	60
GAAAAAGTAG CAGAAAATAA AGAGAAACAT GAAAATATCC ATAGTGCTAT GGAAACTTCA	120
CAGGATTTTA AAGAGAAGAA AACAGCAGTC ATTAAGGAAA AAGAAGTTGT TAGTAAAAAT	180
CCTGTGATAG ACAATAACAC TAGCAATGAA GAAGCAAAAA TCAAAGAAGA AAATTCCAAT	240
AAATCCCAAG GAGATTATAC GGACTCATTT GTGAATAAAA ACACAGAAAA TCCCAAAAAA	300
GAAGATAAAG TTGTCTATAT TGCTGAATTT AAAGATAAAG AATCTGGAGA AAAAGCAATC	360
AAGGAACTAT CCAGTCTTAA GAATACAAAA GTTTTATATA CTTATGATAG AATTTTTAAC	420
GGTAGTGCCA TAGAAACAAC TCCAGATAAC TTGGACAAAA TTAACAAAAT AGAAGGTATT	480
TCATCGGTTG AAAGGGCACA AAAAGTCCAA CCCATGATGA ATCATGCCAG AAAGGAAATT	540
GGAGTTGAGG AAGCTATTGA TTACCTAAG TCTATCAATG CTCCGTTTGG GAAAAATTTT	600
GATGGTAGAG GTATGGTCAT TTCAAATATC GATACTGGAA CAGATTATAG ACATAAGGCT	660
ATGAGAATCG ATGATGATGC CAAAGCCTCA ATGAGATTTA AAAAAGAAGA CTTAAAAGGC	720
ACTGATAAAA ATTATTGGTT GAGTGATAAA ATCCCTCATG CGTTCAATTA TTATAATGGT	780
GGCAAAATCA CTGTAGAAAA ATATGATGAT GGAAGGGATT ATTTTGACCC ACATGGGATG	840
CATATTGCAG GGATTCTTGC TGGAAATGAT ACTGAACAAG ACATCAAAAA CTTTAACGGC	900
ATAGATGGAA TTGCACCTAA TGCACAAATT TTCTCTACA AAATGTATTC TGACGCAGGA	960
TCTGGGTTTG CGGGTGATGA AACAAATGTTT CATGCTATTG AAGATTCTAT CAAACACAAC	1020
GTTGATGTTG TTTTCGGTATC ATCTGGTTTT ACAGGAACAG GTCTTGATAG TGAGAAATAT	1080
TGGCAAGCTA TTCGGGCATT AAGAAAAGCA GGCATTCCAA TGGTTGTCGC TACGGGTAAC	1140
TATGCGACTT CTGCTTCAAG TTCTTCATGG GATTTAGTAG CAAATAATCA TCTGAAAATG	1200
ACCGACACTG GAAATGTAAC ACGAACTGCA GCACATGAAG ATGCGATAGC GGTGCTTCT	1260
GCTAAAAATC AAACAGTTGA GTTTGATAAA GTTAACATAG GTGGAGAAAG TTTTAAATAC	1320
AGAAATATAG GGGCCTTTTT CGATAAGAGT AAAATCACAA CAAATGAAGA TGAACAAAA	1380
GCTCCTAGTA AATTAATAAT TGTATATATA GGCAAGGGC AAGACCAAGA TTTGATAGGT	1440
TTGGATCTTA GGGGCAAAAT TGCAGTAATG GATAGAATTT ATACAAAGGA TTTAAAAAAT	1500
GCTTTTAAAA AAGCTATGGA TAAGGGTGCA CGCGCCATTA TGGTTGTAAA TACTGTAAAT	1560

TACTACAATA GAGATAATTG GACAGAGCTT CCAGCTATGG GATATGAAGC GGATGAAGGT	1620
ACTAAAAGTC AAGTGTTTTC AATTTCAGGA GATGATGGTG TAAAGCTATG GAACATGATT	1680
AATCCTGATA AAAAACTGA AGTCAAAGA AATAATAAAG AAGATTTTAA AGATAAATTG	1740
GAGCAATACT ATCCAATTGA TATGGAAAGT TTTAATTCCA ACAAACCGAA TGTAGGTGAC	1800
GAAAAAGAGA TTGACTTTAA GTTGCACCT GACACAGACA AAGAACTCTA TAAAGAAGAT	1860
ATCATCGTTC CAGCAGGATC TACATCTTGG GGGCCAAGAA TAGATTTACT TTTAAAACCC	1920
GATGTTTCAG CACCTGGTAA AAATATTAATA TCCACGCTTA ATGTTATTAA TGGCAAATCA	1980
ACTTATGGCT ATATGTCAGG AACTAGTATG GCGACTCCAA TCGTGGCAGC TTCTACTGTT	2040
TTGATTAGAC CGAAATTAATA GGAAATGCTT GAAAGACCTG TATTGAAAAA TCTTAAGGGA	2100
GATGACAAAA TAGATCTTAC AAGTCTTACA AAAATTGCC TACAAAATAC TGC GCGACCT	2160
ATGATGGATG CAACTTCTTG GAAAGAAAA AGTCAATACT TTGCATCACC TAGACAACAG	2220
GGAGCAGGCC TAATTAATGT GGCCAATGCT TTGAGAAATG AAGTTGTAGC AACTTTCAAA	2280
AACACTGATT CTAAAGGTTT GGTAAACTCA TATGGTTCCA TTTCTCTTAA AGAAATAAAA	2340
GGTGATAAAA AATACTTTAC AATCAAGCTT CACAATACAT CAAACAGACC TTTGACTTTT	2400
AAAGTTTCAG CATCAGCGAT AACTACAGAT TCTCTAACTG ACAGATTAATA ACTTGATGAA	2460
ACATATAAAG ATGAAAAATC TCCAGATGGT AAGCAAATG TTCCAGAAAT TCACCCAGAA	2520
AAAGTCAAAG GAGCAAATAT CACATTTGAG CATGATACTT TCACTATAGG CGCAAATCT	2580
AGCTTTGATT TGAATGCGGT TATAAATGTT GGAGAGGCCA AAAACAAAA TAAATTTGTA	2640
GAATCATTTA TTCATTTTGA GTCAGTGGAA GCGATGGAAG CTCTAACTC CAGCGGGAAG	2700
AAAATAAACT TCCAACCTTC TTTGTGATG CCTCTAATGG GATTTGCTGG GAATTGGAAC	2760
CACGAACCAA TCCTTGATAA ATGGGCTTGG GAAGAAGGGT CAAGATCAAA AACACTGGGA	2820
GGTTATGATG ATGATGGTAA ACCGAAAATT CCAGGAACCT TAAATAAGGG AATTGGTGGA	2880
GAACATGGTA TAGATAAATT TAATCCAGCA GGAGTTATAC AAAATAGAAA AGATAAAAAAT	2940
ACAACATCCC TGGATCAAAA TCCAGAATTA TTTGCTTTCA ATAACGAAGG GATCAACGCT	3000
CCATCATCAA GTGGTTCTAA GATTGCTAAC ATTTATCCTT TAGATTCAAA TGGAAATCCT	3060
CAAGATGCTC AACTTGAAAG AGGATTAACA CCTTCTCCAC TTGTATTAAAG AAGTGCAGAA	3120
GAAGGATTGA TT	3132

(2) INFORMATION FOR SEQ ID NO: 111:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 14672 base pairs

806

(B) TYPE: nucleic acid
 (C) STRANDEDNESS: double
 (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 111:

CGAGATTTCT TTAAATGAAC TACGTGAAAT CTACCCATCA TCCAGATCTG GATATCTCT	60
CCTATCTATA AGTAAAGTTT TAGGAGATTT TAATATAAGT TCTCATGCTT TTAAAGCTTC	120
GGTAAGAGAT TTAACCACGC TCAGTTTCCC ACTCATTGCG TTCTGGGAGA GTTCTCATT	180
TATTATTCTT GAAAAAATTA GTAAAAACAA GTTTTATATT TTAGATCCTG CAAAAGGCAG	240
GCAGAGAATG TCAATAAGTG AATTGAAAG GCATTATTCA AATATCATTT TAACATTTAA	300
AAAGTTAGAT AGCTTTATGT CTCGTAAAGA TAATAAGAAG TCGCCTGTTT TAAAGTATTT	360
TTTAAAGTAT AGGAATAAGC TAGGGATTTT ATTTTTTGTA ACAGCATTAT TGTATGTAAT	420
ACAATCATTG GTACCTATAG CTAATAGATA CATAATTGAC ACGAATTTCA AGGACGATTC	480
GTATTCGTCT AGAATGTTAT TTACTATATT ATTTATATTT ACTGTTTCAT TCTCCTAAT	540
GTATTTATTA AGACAGATAT ATGTTGCATC CTTAAAATAT ATAATGGATA AAGAGATTAG	600
CTATGATTTT ATGAAACATT TGATATATTT ACCTTACAGT TTTTATGAAA AACGTACTTT	660
AGGGGATATA CTTTTTAGAG CTAACCTCTAT TGTTTATATA AGAGAAATAC TATCAAATAA	720
TTTTATAGCA GCTATACTTG ATTTGTTAAT GATTGTGGTT TATGCTGTGG TTTTATTTAG	780
CTTTTCTAAG TACATGGTAA TCTTTTAAAT ATCACTAAGT CTAGCTCTAT CTATGTAAAT	840
GTATCCAATC ATAAAAATCT CAAAAAATTT AATTGATAAA AATATAAAAG AAAAGGTTAA	900
TGTTCAAAAT ATTACTCCG AAGTAATTTT TAAAAATAGT GATATTAAGC TAACTGGAGA	960
AGAGGAATTT TGGATTAACA AATGGGATAA TTTAATACA AAACAGCTCA TCATAGGTCG	1020
AAAACCTGAT ATACATTTAT CAATTGTTAG TAGTATAACG AATGTTTTAC AAATATTCT	1080
CCCTGTTTTG ACCCTTATG TAGGTGTAAT TATAAAAACA TTCGAACAAT TGACGTTAAG	1140
ACAAATTGTA GCAATAAGTA CAGTCTCACC ATACTTTATT TCTCCTATAA TTTCTTTAAG	1200
TGATAACTAT ATACAATTAA TGTATTAAA GGGATATTTT TTAAGAATAG AGGATGTGTT	1260
TAATACTAAA TCCGAATTAA TTCCAGAAAG AGTCAGTCAA GATATAAAAT TTGATAAAAA	1320
AATAGAATTA AAAGATATTT GGTATAAATA TGGATTATTT GATGATTATG TTTTGAAAGG	1380
AATAAATGTT ACTATTAAAA AAGGAGAAAC TGTGCTATT GTTGGAGAAT CAGGTTGAGG	1440
TAAGAGTACA TTAGCTAAAA TTTTATTAGG TTTATTAGAA CCTAATATTG GTTCAATAGA	1500
AGTTGATGGA GTAGAAAAAG AAGAAATGG TCAACATTG TATAGAAAGA TTTTGGAGC	1560

AGTGTACAA AATTCAACCC TAAGTTATGG TACCTTAAGA GAGAATTGA CATTTGGACA 1620
 CTTTGTTCFA GATGAAGAAT TAATGACAAA TCTAAATCA ATTGGTCTTA GCAATGTAGT 1680
 TAAATCTTTA CCTCTTGGAT TAGAGACAAT CATCGCTGAA GAAGGTAATA ACTTTTCTGG 1740
 AGGGCAGCAG CAAATGATAC TTFTAGCTCG TTGTCTTTTG TCGAAACCTT CGGTAGTTGT 1800
 TTTGGACGAA GCAACAAGTA GTTTAGATAA TTTATCTCAA CAAATTACAA CTTCTTACTT 1860
 AAGTGAATC GGTACCACTA AGATTTTAAT TGCCCATCGA CTAGATACTA TCAAGTCTGC 1920
 AGATAAGATC TTAGTAATGC ATAATGGTGA AATTGTAGAG ATTGGGACCC ATAGAGAACT 1980
 TCTTGAACFA GGAGGCATTT ATAAGCAATT GTATTCAAAT AATTAGTTT TGATTAAAAG 2040
 GGTAATTTA TGAAGATTAT GAAAAAAAAA TATTGGACTT TAGCGATATT ATTCTTTTGT 2100
 TTGTTCAATA ATTCTGTAC TGCTCAAGAA ATACCTAAA ATCTTGATGG CAATATAACT 2160
 CACACTCAGA CTAGCGAAAG TTTTCTGAA TCTGATGAAA AACAGGTGA CTATTCTAAT 2220
 AAAAATCAAG AAGAAGTAGA CAAAAATAA TTTCGTATTC AAATCGATAA GACAGAATTA 2280
 TTTGTAACAA CAGATAAACA TTTAGAAAAA AACTGTGTGA AATTGGAECT TGAACCACAA 2340
 ATAAATAACG ATATTGTAA CTCTGAAAGT AATAATTTAC TAGGCCAAGA TAATTTAGAT 2400
 AATAAAATTA AGGAAAATGT TTCTCATCTA GATAATAGAG GAGGAAATAT AGAGCATGAC 2460
 AAAGATAACT TAGAATCGTC GATTGTAAGA AAATATGAAT GGGATATAGA TAAAGTTACT 2520
 GGTGGAGGCG AAAGTTATAA ATTATATTCT AAAAGTAATT CTAAAGTTTC AATTGCTATT 2580
 TTAGATTCAG GAGTCGATTT ACAAATACT GGATTACTGA AAAATCTTTC AAATCACTCA 2640
 AAAAATATG TCCCAATAA AGGATATTTA GGAAAAGAGG AGGGAGAGGA AGGAATAATA 2700
 TCAGATATTC AAGATAGATT AGGTCAATGG ACGGCTGTTG TAGCTCAAAT TGTAGGGGAT 2760
 GACAATATTA ATGGAGTAAA TCCTCACGTT AATATTAACG TCTATAGAAT ATTTGGTAAG 2820
 TCGTCAGCTA GTCCAGATTG GATTGTAAAA GCAATTTTGT ATGCTGTAGA TGATGGCAAT 2880
 GATATATCA ATCTTAGTAC TGGACAATAT TTAATGATTG ATGGAGAATA TGAGGACGGA 2940
 ACAAATGATT TTGAAACATT TTTGAAGTAT AAAAAGGCTA TTGATTACGC GAATCAAAAA 3000
 GGAGTAATTA TAGTAGCTGC ATTAGGGAAT GACTCCCTAA ATGTATCAAA TCAGTCAGAT 3060
 TTATTGAAAC TTATTAGTTC ACGCAAAAAA GTAAGAAAA CAGGATTAGT AGTTGATGTT 3120
 CCAAGTTATT TCTCATCTAC AATTTCCGTC GGAGGCATAG ATCGCTTAGG TAATTTATCA 3180
 GATTTTAGCA ATAAAGGGGA TTCTGATGCA ATATATGCGC CTGCAGGCTC AACATTATCT 3240
 CTTTCAGAAT TAGGACTTAA TAACTTTATT AATGCAGAAA AATATAAAGA AGATTGGATT 3300

TTTTCGGCAA	CACTAGGAGG	ATATACGTAT	CTTTATGGAA	ACTCATTTC	TGCTCCTAAA	3360
GTTTCTGGTG	CGATTGCAAT	GATTATTGAT	AAATACAAAT	TAAAAGATCA	GCCCTATAAT	3420
TATATGTTTG	TAAAAAAT	CTGGAAGAAA	CATTACCAGT	AAAAATGGT	ATAAAAAGTGT	3480
TAAATATACC	AAACGTATTG	AGATATGATT	TGAATATGTT	ACAATTAGAA	TATAAAAAATG	3540
AACAAAGTTG	GGATAGTTTC	ATAGATAATG	TTAATTTAAT	TGAGTTGGAA	GAGAGAATTC	3600
AAACTACTAT	TGGAATTAAA	CAAATAAACA	CACACAATAT	TATTACTATT	GCCCGAGAAG	3660
GGTACTCTCA	AAATTATTTA	CCTAACACTT	CAGAAAATAC	ATATAATCA	TTACAAGTCA	3720
GTTTAGTTGG	AGTATTACTA	CTTTTTATAA	GTATGGTAAA	TATTTTATGG	GCTAAAAAAA	3780
GTAAATGAAA	ATAAAATTTG	GAGCCCTCTG	AAAAAGTAAG	TCCTACAGTT	CAACTAAAAAT	3840
GAGTCAAAAG	ATGAATCACC	TTGATGTAGG	GGAGTTTGTC	TTATTGCTGC	CTGAACACCT	3900
CCGTTTCAGAG	GAAGAACATT	ATAAATCTGT	TTTGAAGAC	GACTTAACCA	GTCGCATATC	3960
TAGTCAAGAT	GAACGACAGC	AAATGACTGC	TACGGTAGGT	TATTTAGAAT	CAGGTCAGGA	4020
TCGTTTTGTG	TATAATACGA	CCCCTATTTT	TTACCAGCAG	TTTTTGAAAG	ATCCAATCAT	4080
CATTGTATA	ACACCCCAAT	CAACTGGTCC	ACAGTCCATT	TTGTTTTGGA	TAGACGCAGT	4140
ACAGAACTAC	GTTCTCTTTA	ATCAATGTGC	TGATGCCAG	GAGCTTATCC	AGAGACAAGG	4200
CATTGAAAAT	TGGGTCTCAG	AAATGCAAAC	AGGTTACCAC	AACTACATCA	CATTATTGGA	4260
TAATATCCAG	AGGGAACGTT	GGGTAATGCT	AGCAGGAGCT	GTGCTTGGGA	TTGCAACTTC	4320
AATCTTGTG	TTTAACACTA	TGAATAGGCT	CTACTTTGAA	GAATTTAGAC	GTGCCATTTT	4380
TATCAAACGC	ATTGCAGGTC	TCAGGTCTTT	AGAAATCCAT	CGCACTTATC	TCTTTGCTCA	4440
ACTGGGTGTG	TTTTTACTGG	GATTTGTTGC	GAGTGTATTT	CTTCAGGTAG	AGATAGGAGT	4500
TGCTTTCTTA	GTCTTGTTAC	TCTTTACTGG	TCTATCTCTT	TTACAGTTAC	ATGTCCAAAT	4560
GCAGAAAGAA	AACAAGATGT	CCATGCTTGT	TTTGAAGGGA	GGTTAATATG	ATTGAACTTA	4620
AACAGGTGAG	TAAATCTTTT	GGAGAACGAG	AGTTATTTTC	GAATCTTTCA	ATGACATTTG	4680
AGGCTGGAAA	AGTCTATGCC	TTAATTGTTT	CAAGTGGTAG	CGGAAAAACA	ACCTTGATGA	4740
ACATGATTGG	GAAATTAGAA	CCTTATGATG	GGACGATTTT	TTACCGAGGT	AAAGACTTGG	4800
CCAATTATAA	ATCAAGTGAT	TTTTTCCGTC	ACGAATTGGG	CTACCTCTTC	CAGAACTTTG	4860
GCTTAATTGA	AAACCAAAGT	ATTGAAGAAA	ACCTTAAGCT	AGGTCTCATT	GGTCAAAAGT	4920
TGAGTCGGTC	GGAACAGCGG	TTGAGGCAGA	AGCAGGCTTT	AGAACAGGTC	GGCCTGGTTT	4980
ATCTTGACCT	AGATAAGCGC	ATCTTTGAGT	TATCGGCGG	AGAATCGCAA	CGGGTTGCCT	5040
TGGCAAAAAT	TATCTTAAAG	AATCCACCCT	TTATTCTGGC	AGATGAGCCA	ACAGCTTCAA	5100

809

TAGACCCAGC AACCTCTCAG TTGATTATGG AGATTTTGCT ATCTCTTCGA GATGATAATA	5160
GGCTAATCAT TATCGCAACA CATAATCCGG CAATTTGGGA GATGGCTGAT GAAGTGTTC	5220
CGATGGATCA TCTGAAATAA AAATCCTTGT TTTAATTGC ACGATGAGTT ACTGAAATAT	5280
TATCATGAAT CAAGAATTGG AGTTAATTTA GAATGTACT TAATTTAGAA TTGTACTTTA	5340
TTAATATTGA GGTAACTTT TCTTGATAAA GGAAGAAATA ATGGAGAGGA AGTTAGAATG	5400
AAAAAATTCG ACAATTATAT TATTGAGAAG CCTTGCGATT CTAATTCAGA TAAACTGCAA	5460
AAAATCTTAA TAATTGAAAG TTGGTAGAT GATATTTTGC AATTTTCTCT CAGAATCAAT	5520
AATAGTGTAG GAGAGATTTT CCTCCTACAA CCGTTTTAAA AGAAAACAT CTTTATTCCA	5580
TGTTATTTTG AGGAAGATAT TGTGAAAGTC AAAGATGATG ATAAAGTTGA GTGGAATTTG	5640
TTAGAATTC AAAAATTTAG AGCATTTTTG GCTTAGTAAT CTGTGTTGAA GGCTCAAAAC	5700
CTATGGTAAA AAAGTAGCTT TGAACACGTA TTGCCTCCAA AGATTTAGTT AAATAATGAT	5760
TTAACACAAA AAGAAATAT TGAAGTCTG GAAAGATGTT GTTTCAGTAT TGAGAAAAGG	5820
TGGGAAAAAC TTGCGATTTT CACAGAGAAA GGAAGAAAAA GTATAGAAAT ATAGTCAATT	5880
GAAACAAGAA CAGGATAAAA GAACCTTTTG TGCCATATTT TTCTCCTTTC GCTTTACAAT	5940
TGGATTGAAC ACCTTTATTG TATCGCGTTT GGAGTTTTTT TGGTATAACC TTCGACGCAC	6000
ACCCGCATAG CGGGTGTTTT TTTTGTCTCG CACCTAACGG AGCGAGACAA ACTAATAGTC	6060
ACTTAATCAA AAAACGCACC ATATCAAAAA CTAAAAAGTT TGATATCATG CGTCATGTCT	6120
TAAACTAATT GACTATACTT TCTATTCAA TGAGCTTTTA ACCAATGAT TGAGCCAATC	6180
CACTCTFAAA ACCAAAGAGC AATTTCTCGC TTAGCTGACT CTCTGAATC TGAACCATGT	6240
ACAACATTTT GGATAATCTC ATTTTCTCCA GCAGCTTTTG CAAAATCACC TCGAATAGTG	6300
CCTGGTAAAG CTCTTCTGG ACGAGTTGCA CCCATCATGG TCCGCCAAGT TTCGATTACT	6360
TTGGGACCAG AAATGACACC CACAAGAACT GGACCTGAAG TCATGAATTC ACGAATCGGT	6420
GGGTAAAAAC TCTGACCAAC CAAGTCCTGA TAGTGCTGGT CAATCAACTC TTCTGAAACC	6480
TGTGAACGAA ACTCCAATTT TFCGATTGTA AATCCACGTT GTTCGATGCG CTTAACACT	6540
TCACCCACTA GCCCTCTTTT TACACCATCT GGTTTGATGA TAAAGAATGT TTGTTCCATA	6600
CCCGTCTCCT TTGTGAGCTT CTTTCTTTTA TTTTACCACA TTTCGTGGAA AAATGGAGAA	6660
AGTTTTCAGA AGAGAGAATG AGAGAACCCT CGGGTCTCTT CATTCTCTCT TATTCTACTG	6720
TTTCTTCCAC AGTTTCAACG GCAGTATCCA CAACTACTTC TGTTGTTTCT TCATTTCTTT	6780
CTTCCTCTAC TGGAGGATTA AGGTATTCTT CTTCGTTGAC AGCATGTGGT TCAAGGTTAC	6840

GGTAACGGGC	CATACCAGTA	CCAGCTGGGA	TGATCTTACC	GATGATAACA	TTTTCTTTAA	6900
GTCCAAGGAG	ATGGTCTTTC	TTACCACGGA	TAGCTGCGTC	AGTAAGGACA	CGAGTTGTTT	6960
CCTGGAAGGA	AGCCGCTGAC	AAGAACTGT	TTGTTTCAAG	TGAGGCTTTG	GTAATTCCCA	7020
TAAGGACTGG	GCGACCTGTC	GCTGGAATC	CACCTGCGAT	AAGGACATCT	TTGTTGGCAT	7080
CTGTAAAGTC	ATTGATATCC	ATGAGGGTAC	CCATGAGAAG	ATCTGTATCA	CCTGGATCCA	7140
TGACACGGAC	TTTACGGATC	ATTTGACGAA	CCATTACCTC	GATGTGTTTG	TCACCGATTT	7200
CTACCCCTTG	GCTACGGTAA	ACTTTTTGTA	CTTACCAGAG	AAGGTACGTT	TCAACTGACA	7260
AGACATCAG	AACTGCAAGG	AGACGTTTTG	GTTGGATAGA	ACCTTCTGTC	AGAGCAGCAC	7320
CACGCGTAC	TTGGCCCCA	ACTTCGACAC	GCATACGAGC	TGTAATGGA	ACGACATATT	7380
CACCTTCGCC	AGTTTCACCC	TTAACAAAGA	CTTCTTGGT	ACGAGTTGAT	GCATCTTCTT	7440
CGATAGCAGT	AACTTGTCCT	TTAACCTCTG	TAATAACCGC	TTCCCCTTA	GGATTGCGGG	7500
CTTCAAAGAT	TTCTTGACA	CGAGGAAGAC	CCTGAGTGAT	ATCGGTATTT	GAGGCAACCC	7560
CACCTGTGTG	GAAGGTACGC	ATTGTAAGCT	GTGTACCAGG	TTCCCCGATA	GATTGGGCAG	7620
CGATTGTACC	AACTGCTTCA	CCAACCTCAA	CCGCATCACC	AGTCGCCAAG	TTGATACCGT	7680
AACAGTGACG	GCAGACACCG	TGACGAGTGT	TACATGTAAA	TACAGAACGG	ATAGTCACTT	7740
CTTCCACACC	AGCATGACA	ATTTACGCG	CCTTGCTTTC	TGTAATCAAT	TCATTTGGAC	7800
CAATAATCAC	TGCACCAGTT	TCTGGATGTT	TAACAGTTTT	CTTAGTGTA	CGACCGTTGA	7860
GACGCTCTTC	GAGAGACTCG	ATCATCTCTT	TTCTTCTGC	GATAGAACGG	ATCAAGAGAC	7920
CACGGTCAGT	TCCACAGTCG	TCCTCACGGA	TGATAACGTC	TTGGGCAACG	TCGACCAAAC	7980
GACGAGTCAA	GTAACCTGAG	TCGGCTGTCT	TAAGGGCCGT	ATCGGTCATA	CCTTTACGAG	8040
CACCGTGAGT	TGAGAAGAAC	ATTTCCAATA	CCGACAAACC	TTCGCGGAAG	TTTGAAAGGA	8100
TTGGCAATTC	CATGATACGT	CCATTCGGAG	CAGCCATCAG	ACCACGCATA	CCGGCAAGCT	8160
GTGAGAAGTT	TGAGATGTTA	CCACGGGCTC	CAGAGTCCAT	CATCATAACG	ATTGGGTTCCT	8220
TAGGATCTTG	GTTAGCAATC	AAGCGTTTCT	CAAGTTTTTC	ACGGGCAGCA	CGCCATTCAG	8280
CTGTAAACAGC	ATTGTAAACG	TCGTCGTCTG	TGATCATACC	ACGACGGAAT	TGTTTGGTGA	8340
TTTGTTCGAC	ACGTTTGTGT	GATTCTTCAA	TGATTTCAGC	CTTGTATCA	ACGACTGGGA	8400
TATCGGCAAT	ACCCACTGTC	AATCCTGCAA	GAGTTGAGTG	GTGGTAACCG	AGGTTCTTCA	8460
TGCGGTCAAG	TAGGGCAGAA	GTTTCTGTCTG	TACGGAAACG	TTTGAAGATT	TCAGCGATGA	8520
TATTTCCAAG	GTTTTTCTTC	TTGAATGGAG	GGTTGAGCTC	AAGATGCTG	ATAGCTTCCT	8580
TGATATCTCC	ACCAAGTGGC	AAGAAGTATT	TAGCTGGAAC	ACCTTCTGTC	AAGTTGGCAT	8640

TGTTTGGTTC	TTGCAAGTAT	GGTAGCCCCT	CTGGCATGAT	ATCGTTGAAG	AGAATTTTAC	8700
CAACTGTTGT	AAGCAAGACC	TTATGTCTTT	GCTCTTCTGT	CCAAGGCTTG	TTGAGGCTGT	8760
CTGTGTCGAT	ACCAACACGT	GAGTGGAGGT	GAACATAACC	ATTGCGGTAA	GCCATAACCG	8820
CTTCGTCACG	GTCTTTGAAG	ACCATTCCCT	CACCTTCGCG	ACCAGCTTCT	TCCATGGTCA	8880
AGTAGTAGTT	ACCCAAAACC	ATGTCCTGAG	ATGGAGTAAC	TACCGGTTTC	CCATCTTTCG	8940
GGTTCAAGAT	GTGCTCAGCA	GCTAGCATGA	GGATACGAGC	TTCTGCTTGT	GCTTCTTCTG	9000
AAAGTGGTAC	GTGGATGGCC	ATTTGGTCCC	CGTCAAAGTC	AGCATGTAG	GCTTCACAGA	9060
CAAGTGGGTG	CAAGCGAAGA	GCCTTACCAT	CAATCAAGAC	TGGCTCGAAG	GCTTGGATAC	9120
CCAAACGGTG	AAGGTCGGT	GCGCGGTTCA	AAAGCACTGG	GTGTCTTTA	ATCACTTCTT	9180
CAAGGATATC	CCAGATACGC	TCATCTCCGC	GTTCCACCAA	GCGTTTAGCT	GCTTTGACGT	9240
TTTGACGAT	ATCACGGCA	ACGATTTCAC	GCATGACAAA	TGGTTTAAAG	AGTTCAATCG	9300
CCATTTACG	CGGCACACCA	CATTGGTACA	TCTTAAGAGT	TGGACCAACG	GCGATAACTG	9360
AACGTCCTGA	GAAGTCAACA	CGTTTACCGA	GCAAGTTTGG	ACGGAAGCGT	CCTTGTTTAC	9420
CTTTAAGCAT	GTGGCTCAAT	GATTTCAATG	GACGGCTACC	TGGTCCTGTG	ATTGGACGAC	9480
CACGACGACC	ATTGTCAATC	AAAGCGTCAA	CTGCTTCTTG	AAGCATACGC	TTCTCATTTT	9540
GAACGATGAT	ACCTGGTGCA	TTTAACTCAA	GCAAACGAGC	CAAACGGTTG	TTACGGTTGA	9600
TAACACGGCG	GTAAAGGTCA	TTCAAGTCAG	ATGAGGCAAA	ACGGCCACCA	TCCAAGTCA	9660
ACATTGGACG	AAGATCTGGT	GGGATAACCG	GAAGATGTT	AAGAATCATC	CATTCAGGTT	9720
TGTTTCCAGA	CTTGTA AAAAG	GCATCCAAAA	CATCCAAACG	ACGGATGGCT	TTGACACGCT	9780
TTTGTCAGT	AGCTGTTTTC	AATTCTTCTT	TGAGTTCAGC	AATTTCTTTT	TCAAGATCTA	9840
CTTGCTTCAA	AAGGTCTTGG	ATGGCTTCCG	CACCCATCTT	GGCAACAAAT	GAACCATAAC	9900
CATATTCACG	CAAGCGCTCT	CGGTATTCGC	GCTCTGTGAT	GATAGACTTG	TGCTCAAGTG	9960
GTGTATCCCT	AGGATCAATC	ACCACATAAG	CCGCAAAGTA	GATAACTTCC	TCGAGGGCAC	10020
GAGGGCTCAT	ATCAAGGGTC	AAGCCATAC	GGCTTGGAAT	CCCCTTGAAG	TACCAGATGT	10080
GAGATACAGG	AGCTTTCAAT	TCGATATGTC	CCATACGCTC	ACGACGAACT	TTCGTACGCG	10140
TTACTTCAAC	CCCACAGCGG	TCACAAACAA	TTCTCTGTGA	ACGAATGCGT	TTGTACTTAC	10200
CACAAGCACA	TTCCAGTCT	TTTGTAGGAC	CAAAGATCAC	TTCATCAAAG	AGTCCTTCAC	10260
GTCTGTTT	CAAGGTACGA	TAATTGATTG	TTTCAGGTTT	TTTGACTTCT	CCATAAGACC	10320
ATGAACGGAC	TTTACTTGA	GAAGCTAGGG	TGATTTGCAT	ACTTTTAAAA	CGATTACAT	10380

CAACCACTAT TTCTTCCCTT TCTATTCTAA GTGAACCTGCT TATPCTTGTT CAGCAGCTTC 10440
 TTCTGTTGCT TCCGCTTTTG TTGCTTTCTC AGCTTCTTCA GCTTCAAAGG CTGCTTTAGC 10500
 CTCTTGGGCT GCTTTTTTCGC GGGCTTTTTC AAGGTCATCT ACGTGGATGA CATCTTCGTC 10560
 CATTCCCTCA TCCAAGTCGC GAAGTTCAC TTCTTGGTCA TCTTCGTCTA GGACACGCAT 10620
 GTCAAGACCA AGAGATTGCA ATTCTTTGAC AAGAACCTCGG AAGGATTCTG GAACACCTGG 10680
 TTTTGGAATT GGTTTGCCTT TTGTAATAGC TTCATAGGCT TTCAAACGTC CGTTGATATC 10740
 GTCCGACTTG TAAGTCAAGA TTTCTTGAAG GACATTTGAC GCACCGTAGG CTTCAAGAGC 10800
 CCAAACCTCC ATCTCACCGA AACGTTGTCC ACCAAACTGA GCCTTACCTC CGAGTGGTTG 10860
 TTGGGTAAACA GTTGAGTATG GTCCGACTGA ACGCGCGTGC AATTTATCAT CAACCATGTG 10920
 GTGGAGTTG ATCATGTACA TGACTCCGAC AGAAACACGG TTATCAAACG GTTCACCAGT 10980
 ACGTCCATCG TAAAGGATCG TTTTGGCATC GCTATCCATA CCTGCTTCTT TAACAGTTGA 11040
 CCAAAGATCT TCAGAACTTG CTCATCAAA GACTGGTGTA GCGATGTGAA TACCAAGAGT 11100
 ACGAGCTGCC ATACCAAGGT GAAGCTCCAT AACCTGACCG ATATTCATAC GTGATGGTAC 11160
 CCCAAGTGGG TCAACATGA TGTCGACTGG AGTTCCTCTT GGAAGGTAAG GCATGTCTTC 11220
 TACAGGAACG ATACGAGAGA CAACCCCTTT GTTCCGTGA CGTCCGGCCA TTTTATCTCC 11280
 GACCTTAATC TTACGTTTTT GAGCGATGTA AACACGAACC AACATGTAA CACCTGATTG 11340
 CAACTCATCT CCATTTACAC GTGTAAAGAT CTTAACATCA CGAACGACAC CATCGGCACC 11400
 GTGTGGTACA CGAAGAGAAG TATCACGCAC TTCACGAGAC TTGTCTCAA AGATAGCGTG 11460
 CAAGAGACGT TCTTCAGCTG AAAGATCTTT CTCACCCCTA GGTGTTACTT TACCTACAAG 11520
 AATATCACCT TCTTTAACCT CAGCACCAAT ACGGATAATC CCCATTTTCGT CAAGGTCTTT 11580
 GAGGGCATCT TCACCAACGT TTGGAATTTT GCGAGTGATT TCTTCAGGCC CAAGCTTTGT 11640
 ATCGCGCGTT TCTGATTCGT APTCTTCAAG GTGAACAGAT GTGTAGACAT CGTCCTTCAC 11700
 CAAGCGTTTC CTGATGATAA CGGCATCCTC GAAGTTGTAA CCTTCCAAG TCATGTAGGC 11760
 AACGATTGGG TTTTGTCCAA GCGCCATTTT TCCATTTTCC ATAGAAGGTC CGTCAGCGAT 11820
 GAAATCGCCT TTTTCAACGA CATCACCAAC TTTTACGAGA GTGCGTTGGT TGTAAGCAGT 11880
 ACCTGAGTTT GAACGACGGA ATTTTTGGAT GTGGTAAACA TCCAATGAAC CATCTTCACG 11940
 ACGAACTTCT ACCTTGTCAG CATCTGCGTA AGTAACTTTA CCATCATACT GAGCAATCAC 12000
 AGCCGCACCA GAATCGTGGG CTGCTTGGTA TTCCATACCA GTACCAACGT AAGGTGCCTG 12060
 AGGATTAATC AATGGCACAG CCTGACGTTG CATATTTGGCT CCCATGAGGG CACGGTTGGA 12120
 GTCATCGTTT TCCAAGAAAG GAATACATGC TGTCGCAACG GCAACTACCT GTTTTGGTGA 12180

AACGTCCATG TAGTCAACAA TATTAGCTGG ATACTCTTGG TTGACCCCTT GGTGACGTCC 12240
 CATGACAATC TTCTCAGCAA AGGTTCATC TTCATTGAGA CGAGAGTTAG CCTGAGCTAC 12300
 AGTATATTCA TCTTCTTCAT CAGCTGTCAA CCAAACAATT TCGTTCGTGA CAACACCTGT 12360
 TTCACGGTCA ACCTTACGGT ATGGTGTGTT AACAAAACCA TATTTGTTCA AGTGTCATA 12420
 AGATGACAAG TTATTGATCA AACCGATGTT AGGTCCTTCA GGTGTCTCGA TTGGACACAT 12480
 ACGACCATAG TGAGTGTAGT GCACGTACG TACTTCATAT CCAGCACGGT CACGAGTCAA 12540
 ACCACCAGGT CCTAAGGCTG ACAAACGGCG TTTGTGAGAC AACTCAGAAA GCGGGTTGTG 12600
 TTGGTCCATG AACTGTGACA ACTGTGATGA ACCAAAGAAT TCTTTAACTG CAGCTGTTAC 12660
 AGGACGGATA TTGATAATTT GTTGTGGTGT CAAGACTTCA TTGTCCTGAA CAGACATACC 12720
 TTCACGGACA TTACGTCCA TACGAGAAAG TCCCAAACGT ACTTGGTTGG CAAGCAATTC 12780
 ACCAACCGCA CGGATACGAC GATTTCCAAG GTGGTCGATA TCACTACAC GGCCAAGTCC 12840
 TTCAGCCAAG TTGAGGAAGT AGCTCATCTC AGCAAGGATA TCTGCAGGAG TCACCGTACG 12900
 AACCTTGTC TCTGGGTTAG CATTACCAAT GATCGTTACG ACGCGATCTG GATCAGTTGG 12960
 AGCAATAACC TTGAATTTTT GAAGAACAAC AGGCTCAGTC ACAACGGCTG CATCGTTTGG 13020
 GATGTAGACA ATCTTGTTCA AGTCGCCATC CAAATGGCTT TCAATGCTTT CAATCACGCT 13080
 ACGAGTCATA ATCGTACCAG CTTCTACCAA GATTTCTCCA GTTTCAGGGT CTACCAATGG 13140
 CTCTGCAATG GTTTGGTTGA GCAAACGTGT TTTAACATTG AGTTTTTTAT TGATTTTGTA 13200
 ACGACCAACT GCTGCCAAGT CATAACGACG TGGGTCAAAG AAGCGAGCTA CAAGCAAGCT 13260
 ACGTGAGCTT TCAGCCGTCT TAGGCTCACC TGGACGAAGG CGTTCGTAAA TTTCTTTCAA 13320
 GGCTTCGTCT GTACGAGAGT CCATTGGATT CTTGTGGATA TCTTTTCAA CAGTGTGCG 13380
 AACCAATTCG CTGTCACCAA AGATATCAAA GATTTATCA TCACCTGAGA AACCAAGAGC 13440
 ACGAACCAAG GTTGTAATG GAATCTTACG AGTACGGTCG ATACGAGTGT AGGTGATATC 13500
 TTTTGAGTCG CTTTCAAGTT CCAACCAAGC TCCACGGTTA GGGATAACAG TTGAACCATA 13560
 GCCCACCTTA CCATTTTTGT CTACTTTGTC GTTAAAGTAA ACACCTGGTG AGCGGACCAA 13620
 CTGAGAAACG ATAATACGTT CACCACCATT GATGATGAAA GTACCCATTT CTGTCATGAT 13680
 TGGGAAATCA CCAAAGAAAA CTTCTGGGT CTTGATTTG CTTGTTTCTT TATTGATCAA 13740
 ACGGAAGGTT AAAAAATG GTGCTGAGTA GCTAGCATCG TGGATACGAG CTTCTTCTAG 13800
 CGTATATTTT GGTTCCTTGA TTTCATATCC AACAAATCC AACTCCATG TGTCTGTGAA 13860
 GTTTGAAATT GGCAATACAT CTTCAAACAC TTCCTTAAGA CCGTGGTCTA GGAAAGCTTT 13920

814

GAATGAGTCA GTTTGAATTT CAATCAAATT TGGTAAGTCA AGAACTTCTT TGATTCTTGA 13980
 AAAACTACGA CGGGTACGAT GTTTCCTCGTA TTGAACGTCA TGTCCTGCCA AGATGATTCT 14040
 CCTTTGTAAA TAAGTTCCAA GCCTTGTCAA TCAGGCTTTT CTAATCGTCA TATGGTTGTA 14100
 AACCCCTTAT CACCGTGTCC TCTTGACGAA TTTTCAGAAT CTTTAAGCCT CTGTTACAAA 14160
 TGCTCAAAAT CTTGAAAAAA AGCACAAAAA GAGCAGCTAA ATCTGACTTT TTCAGAAGAT 14220
 TTAACCTGCTG TGAGCCTTGT CTGGACAATA TTTCAGACAA AACCTACGAC AAATGATTAC 14280
 CCATATTATA CCCTATTTAG CTAGATTTTT CAAGGGGTTT CAGTAGGTTT TTGGTAAATT 14340
 TTTTCCATA GAAAACCTGG CATCACATTC GAATCACGCT ATGGTACAAA AAACTGAAAA 14400
 AACTATTGAC TGAAAATCAT TTTCAAGGTA TAATAATAAA CGTTAAGGCG GTATAGCCAA 14460
 GTGGTAAGGC ACGGCTCTGC AAAAGCTTGA TCGTCGGTTC AAATCCGTCT ACCGCCTTCT 14520
 ATAACCTGAT TTATCAGGT TCAAATGAAC AGAAAGCCCA ATTTGAAGGG CTTTTTTTAT 14580
 TTTCCCTCGA ATAAATACGT ATAACTTTAA AACTTTTGG AGCGAGTTTG TGGCAGAGTT 14640
 CTTTCCATGG CATAATCCC TTTTGAAATC AG 14672

(2) INFORMATION FOR SEQ ID NO: 112:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 7902 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 112:

AGGAGACTAT TCAAGCCCAA ATTgAGTAGC CCAGCAAAGA CTGTATAGAC TGTGATACGT 60
 TTTTCATAGC CATTTGGTAAA GAGAATTTGG GAACCAAGAA TGGTATCTAA GGCCAGGATA 120
 ATCGTACGAA AAGCGAAGAG AGAGGTCAAG ATGCCGCCTC CGATATATTT TTCACTACCG 180
 TAAAGTAGGA TGGCATTGG TCCTAAAACC ATGAGTCCAA AACTCAGTGG AATGATAAAG 240
 AAGTTAAAGA TTCGACTACC TCTATTAACC AGAGAAACAT AGGCTTCTTT GTCTCCTTTC 300
 CCCAGATAGT AACTGAGACG AGGCACACTC ACTCCAATTG CACCTGTTAC AACCCAGCT 360
 ATAACGGTCA CAATTCGCTG AGCTATGGTA TAGTAACTAA CGTTGACATC AATCCCTGTT 420
 TTAACGAGGA AGAGGCGATC TAAAAAAGTG AAGAGCATAT TGGCATTGGC AAAGACTAAC 480
 ATGGCTGTCA GAGGGAGAAA GAGTGGTTTA AAATCACTTA GGTGAATTTT AACAAGTTTG 540
 ATGTCTCTTT TAATCCAAAA ATAACTAATC AGGTAGTTAA TCAGCGTCGA TAAACTCATC 600
 ACAAGTGTAT AGACAACAAT ATCGTGTTC TTTTAAACAA ATAAGAAAAT AGAGACCAGC 660

ATCAGGATAC GGATGAAGGC AGTTTTGTAA AAGAGAAAAC TGTAATTTTC CAGAGCTTCA 720
 TTGACCCATT CGATTGAAA AATCTGGGCA ATGAGTTGAA TCCCCATAAC AAGGTAGACC 780
 TTTTGTACGA TTGGATTATC AGTAAAGAAG AGAGGATAGG CTAGGATATA GACAGCAGTG 840
 GTCAAAATCG TACAAGCGAT GCACAAATAA AAAAGACTAG AAAAGGTCT GTTAAGATCT 900
 TTTTGTATTAT CCTTGACATT ACTGATAGCC CTTAAACCGT AGTTATAGAC ACCATAAGTT 960
 GCAAAGGGCA AGAAAAATGA CAAAATAGTG TCGACTGAGT TGAAGTAACC ATAGTCAGTT 1020
 CGGTCCAAGA CACGCGCGAC ATAGGTTCCA GTTAGGATGG GAAAAATAAT ATTCAAGACA 1080
 CGAATCCCA TGTAAGATAG AGCATTTAAT TTTATACTTT TCATTCAATT TACCTCGTTT 1140
 TTCATTATAT CATAAAGTTA GCTAATAAGA AATGAAGGGC AGTAAGTCAA GTAATCACTT 1200
 TGAAGTTTCA AATCTTAAGT TTTAAGTTTT CTTAAGGAA AGTATATTAT TCTGAAGGAC 1260
 TCTAAAATTT CGCAGCCATT TATTAGTAAT TGCTACAGAA TTCCTAGTCA TTAGTAGAAA 1320
 TGGACTAGTT TCTTTGAATA ATAGAAGTGC ATAATCTCC TATTCTAGAA GGGGAGGACC 1380
 AGTATTTCTT TTATGATAGG ACTAGATTGT GGTATAATAG AGAGAATAAG TTTTTTAGT 1440
 AAGACAAAGG AGAAAATAGA TGATTTATGC AGGAATTCTT GCCGGTGGAA CTGGCACACG 1500
 CATGGGGATC AGTAACTTGC CAAAACAATT TTTAGAGCTA GGTGATCGAC CTATTTTGAT 1560
 TCATACAATT GAAAAATTG TCTTGAGGCC AAGTATTGAA AAAATTGTAG TTGGTGTTC A 1620
 TGGAGACTGG GTTCTCATG CAGAAGATCT TGTAGATAAA TATCTTCCCTC TTTATAAGGA 1680
 ACGTATCATC ATTACAAAGG GTGGTGCTGA CCGCAATACA AGTATTAAGA ACATCATTGA 1740
 AGCCATGAT GCTTATCGTC CGCTTACTCC AGAGGATATC GTTGTACCC ACGATTCTGT 1800
 TCGTCCATTT ATTACACTTC GCATGATTCA GGACAATATC CAACTTGCCC AAAATCATGA 1860
 CGCAGTGGAC ACAGTGGTAG AAGCGGTGA TACTATCGTT GAAAGTACCA ATGGTCAATT 1920
 TATTACAGAT ATTCCAAATC GTGCTCACCT TTATCAAGGA CAAACACCTC AAACATTCCG 1980
 TTGCAAGGAC TTCATGGACC TTTATGGATC TCTTCTGAT GAAGAGAAGG AAATCTTGAC 2040
 AGATGCATGT AAAATCTTTG TGATCAAAGG AAAAGATGTG GCTTTGGCCA AAGGTGAATA 2100
 CTCAAATCTG AAGATTACAA CCGTAACAGA TTTGAAGATT GCAAAAAGTA TGATTGAGAA 2160
 AGACTAGTAA AATGATTAAT CAAATTTATC AACTAACTAA GCCTAAGTTT ATCAATGTCA 2220
 AATATCAGGA AGAGGCTATT GACCAAGAGA ATCATATCCT TATCCGTCCC AACTACATGG 2280
 CTGTCTGTCA TGCAGATCAG CGTTACTATC AGGGAAAACG TGATCCCAAG ATTTTGAATA 2340
 AAAAGCTTCC AATGGCAATG ATTCACGAGT CATGTGGAAC CGTCATTTCT GACCCGACCG 2400

GAACCTACGA	GGTTGGTCAA	AAAGTTGTCA	TGATTCCCAA	TCAGTCTCCT	ATGCAGAGTG	2460
ATGAAGAATT	CTATGAAAAC	TACATGACAG	GGACCCATTT	CTTGTCTAGT	GGATTGATG	2520
GCTTTATGAG	AGAGTTTGT	TCTCTCCCTA	AAGATCGTGT	GGTGGCTTAT	GATGCTATTG	2580
AAGATACGGT	TGCAGCCATT	ACAGAGTTTG	TCAGTGTGGG	CATGCACGCT	ATGAATCGTC	2640
TATTGACTCT	TGCTCATAGC	AAGCGGGAGC	GGATCGCCGT	TATTGGAGAT	GGAAGTTTAG	2700
CTTTTGTGGT	TGCCAATATT	ATCAACTATA	CTTTGCCAGA	AGCAGAGATT	GTGGTTATTG	2760
GTGTCATTG	GGAAAAGTTG	GAACTCTTCT	CATTTGCCAA	AGAATGCTAT	ATTACGGATA	2820
ATATTCCTGA	AGATTGGCC	TTTGACCATG	CTTTTGAATG	TTGTGGTGGT	GATGGTACTG	2880
GACCAGCTAT	TAATGACTTG	ATTCGCTACA	TTCGTCTCA	GGGAACGATT	CTCATGATGG	2940
GAGTTAGCGA	ATATAAAGTC	AATCTCAATA	CTCGGATGC	CTTAGAAAAG	GGCTTGATTT	3000
TGGTTGGGTC	ATCTCGTTCT	GGTCGCATTG	ATTTTGA AAA	TGCTATCCAA	ATGATGGAAG	3060
TCAAGAAAT	TGCCAATCGT	CTTAAAAATA	TCCTTTATCT	AGAAGAACCT	GTAAGAGAAA	3120
TTAAAGATAT	TCATCGTGTC	TTTGCAACCG	ATTTAAACAC	AGCCTTTAAA	ACAGTGTTTA	3180
AGTGGGAAGT	ATAAGTACTG	GAGGTTAATT	GTGGAGAAAA	TCATTAAAGA	AAAAATTTCT	3240
TCCTTACTTA	GTCAAGAAGA	GGAAGTCCTC	AGTGTGAAC	AACTGGGTGG	AATGACCAAT	3300
CAAAACTATT	TGGCCAAAAC	AACAAATAAG	CAATACATTG	TTAAATTTCT	TGGTAAAGGG	3360
ACAGAAAAGC	TTATCAATCG	ACAAGATGAA	AAGTACAATC	TTGAACTACT	AAAGGATTTA	3420
GGCTTAGATG	TAAAAAATTA	TCTTTTGGAT	ATTGAAGCTG	GTATCAAAGT	AAATGAGTAT	3480
ATCGAATCTG	CGATTACGCT	TGATCAACG	TCAATCAAGA	CCAAGTTCGA	CAAAATTACT	3540
CCAATATTAC	AAACTATTCA	TACGTCTGCT	AAGGAATTA	GAGGAGAATT	TGCTCCTTTT	3600
GAAGAAATCA	AAAAATACGA	ATCCTTGATT	GAAGAACAAA	TTCCTTATGC	CAACTATGAA	3660
TCTGTTAGAA	ATGCAGTCTT	CTCCTTAGAG	AAAAGACTGG	CTGACTTAGG	TGTTGACAGA	3720
AAATCTTGTC	ATATCGATTT	GGTGCCTGAA	AACTTTATCG	AATCACCTCA	AGGACGACTT	3780
TATTTGATTG	ACTGGGAATA	TTCATCAATG	AATGATCCAA	TGTGGGATTT	GGCTGCCCTC	3840
TTTTTAGAGT	CTGAATTCAC	TTCCCAAGAG	GAAGAACTT	TCTTATCTCA	CTATGAGAGT	3900
GACCAAACAC	CGGTTTCTCA	TGAAAAGATT	GCTATTTATA	AAATTTTACA	AGATACTATT	3960
TGGAGTCTAT	GGACTGTCTA	TAAGGAAGAG	CAAGTGAAG	ATTTTGGTGA	CTATGGTGTG	4020
AATCGTTACC	AAAGAGCTAT	TAAAGGTTTG	GCTTCTTATG	GAGGTTTACA	TGAAAAGTAA	4080
AAACGGAGTT	CCTTTTGGCC	TTCTCTCAGG	TATTTTCTGG	GGCTTGGGTC	TAACGGTTAG	4140
TGCTTATATC	TTTTCGATTT	TTACAGATTT	GTCACCCTTT	GTGGTGGCTG	CAACTCATGA	4200

TTTTTTGAGC ATCTTTATCT TACTAGCTTT TCTCTTGGTA AAAGAAGGGA AAGTTCGCCT	4260
CTCAATTTTC TTAAATATTC GCAATGTCAG TGTATCATC GGAGCCTGTC TAGCAGGCC	4320
TATCGGTATG CAGGCCAATC TTTATGCAGT TAAGTATATC GGAAGTCTT TAGCTTCATC	4380
TGTATCGGCT ATTTACCCTG CGATTTCACT TCTATTGGCT TTCTTCTTTT TGAAGCACAA	4440
GATTTGCGAA AATACTGTAT TTGGGATTGT CTTGATTATT GGAGGGATTA TTGCTCAGAC	4500
CTATAAGGTT GAACAGGTTA ATTCTTTCTA CATTGGGATT CTTTGTGCTT TGGTTTGTGC	4560
TATTGCATGG GGAAGTGAGA GTGTTCTTAG CTCTTTTGCC ATGGAAAGTG AATTGAGTGA	4620
AATCGAAGCC CTCTTAATCC GTCAAGTAAC TTCGTTCTTG TCCTATCTTG TGATTGTGCT	4680
CTTCTCTCAT CAGTCATTTA CTGCAGTAGC CAATGGACAA TTGCTAGGTC TCATGATTGT	4740
TTTTGCAGCC TTTGATATGA TTTCCCTACTT GGCTTATTAT ATCGCTATCA ATCGCTTGCA	4800
ACCAGCCAAG GCTACAGGCT TGAACGTGAG CTATGTAGTA TGGACGGTCT TGTTTGCAGT	4860
TGTTTTCTTG GGTGCACCGC TAGATATGCT GACCATTATG ACGTCACTTG TCGTCATTGC	4920
TGGAGTTTAT ATTATTATTA AAGAATAAAG GAGATTCGTG TGAAAGCCAT TATCTTAGCA	4980
GCGGGATTGG GAACTCGCTT GCGTCCTATG ACTGAAAATA CCCCTAAAGC CTTGGTTCAG	5040
GTTAATCAAA AACCTTTGAT TGAGTACCAA ATTGAGTTTC TCAAAGAAAA AGGAATCAAT	5100
GACATCATCA TCATTGTTGG TTATCTTAAA GAACAATTCG ATTACTTGAA AGAGAAATAC	5160
GGTGTTCGTC TCGTTTTCAA TGATAAATAC GCTGACTACA ATAACTTTTA CTCTCTCTAT	5220
CTTGTA AAAAG AAGAAATTGGC CAACAGCTAT GTTATTGATG CTGACAATTA TCTCTTTAAA	5280
AATATGTTCC GCAATGATTT GACACGTTTC ACTTATTTTA GTGTTTATCG TGAAGATTGT	5340
ACCAACGAAT GGTTCCTGGT TTATGGAGAT GACTACAAGG TTCAAGACAT TATTGTGTGAT	5400
AGCAAGGCAG GTCGCATCCT TAGTGGTGTA TCCTTCTGGG ATGCTCCAAC TGCAGAAAAG	5460
ATTGTCAGCT TTATCGACAA GGCTTATGTA AGTGGTGAAT TTGTTGATCT CTATTGGGAC	5520
AATATGGTTA AGGATAATAT CAAAGAGCTA GATGTCTATG TTGAAGAATT AGAAGGCAAT	5580
AGCATTTATG AGATCGATAG TGTCCAAGAC TATCGTAAAT TAGAAGAAAT TCTTAAAAAC	5640
GAAAATTA AA GATTCCAACA TCTGACAAAA TAGTCGGATG TTTTTTGATT TTTTACGAAC	5700
TTTTACGAAT AGATAGATGA GTAGAAAAAG AAATGGAGTT ATTTATGAAA ATCACAAACT	5760
ATGAAATCTA TAAGTTAAAA AAATCAGGTT TGACCAATCA ACAGATTTTG AAAGTGCTAG	5820
AATACGGTGA AAATGTTGAT CAGGAGCTTT TGTGGGTGA TATTGCAGAT ATCTCAGGTT	5880
GCCGTAATCC AGCCGTTTTT ATGGAACGTT ATTTTCAGAT AGACGATGCG CATTGTGCGA	5940

818

AAGAGTTTCA	AAAATTTCCA	TCTTTCTCTA	TTTTAGATGA	CTGTTATCCT	TGGGATTTGA	6000
GTGAAATATA	TGATGCGCCT	GTACTTTTAT	TTTACAAGGG	AAATCTTGAC	CTCCTGAAAT	6060
TCCCGAAGGT	AGCGGTCGTG	GGCAGTCGTG	CTTGTAGCAA	ACAGGGAGCT	AAGTCAGTTG	6120
AAAAAGTCAT	TCAAGGCTTG	GAAAATGAAC	TGGTTATTGT	CAGTGGTCTG	GCCAAGGGCA	6180
TTGACACAGC	AGCTCATATG	GCAGCTCTTC	AGAATGGCGG	AAAAACCATT	GCAGTGATTG	6240
GAACAGGACT	GGATGTGTTT	TATCCTAAAG	CCAATAAACG	CTTGCAAGAC	TACATCGGCA	6300
ATGACCATCT	GGTCTAAGT	GAATATGGAC	CTGGTGAACA	ACCTCTGAAA	TTTCATTTTC	6360
CTGCCCGTAA	TCGCATCATT	GCTGGACTTT	GTCGTGGTGT	GATTGTAGCA	GAGGCTAAGA	6420
TGCGTTCAGG	TAGTCTCATT	ACGTGTGAGC	GAGCAATGGA	AGAAGGACGC	GATGTCTTTG	6480
CTATTCCTGG	TAGCATTTTA	GATGGACTAT	CAGACGGTTG	CCATCATTTG	ATTCAAGAAG	6540
GAGCAAAAT	GGTCACCAGT	GGGCAAGATG	TTCTTGCGGA	ATTTGAATTT	TAAAAATGAC	6600
CTAAGCTAGA	ATTCTAAGAA	AAAATCAATT	TTAAGAGAAA	ATGAACCCAA	CATTTCCATA	6660
ATAAACCGCA	TATTAGCAAG	TTTTTAACAC	TTGATAATAT	GCGTTTTTTC	TAAGTGGATT	6720
AGTAGAGTAG	AGGATTTTTC	TCATATAATA	CTCTTCGAAA	ATCTCTTCAA	ACTACGTCAG	6780
CTTCCATCTG	CAACCTCAAA	ACAGTATTTT	GAGCgacttc	GTCAGTCTTA	TCTACAACCT	6840
CAAAGCAGTG	CTTTGAGCAA	CCTGTGGCTA	GCTTCCTAGT	TTGCGCTTTG	ATTTTCATTG	6900
AGTATAAGGG	AAAGTATAGT	GAATTGAAAT	AAGATGTGAA	CAACTCTATC	AGGAAAGTCA	6960
AATTAATTTA	TAGAAATATT	TTAGCAGCCA	AGGTGTACTG	TTATAGATTC	AATTACACTA	7020
TAATTTAGTG	TAATTGAGAA	AGGAGAAATG	ATTGTGATTG	ATGTTGGCTA	GGTTATGTTT	7080
AATGATTCCT	ACCGTCTCAA	ATCTTGTCAG	TAAGGAAAAA	TAAATTCTTC	AAAAGTAGAG	7140
ATTACAAGGC	TTGTTTAAGA	AAGAATTCAA	AGACCTTGAC	AAATAAAAAT	AAAATGGTTA	7200
TTATAAAAAA	TGGTCTGAAA	TAGATGATGA	TACTTTTCGA	AAATCTCTTC	AAATACGTCA	7260
GCTCAGCTTT	GCCTTGCTGT	GTTTTGAGCA	AGCTACGGTT	AGCTTCCGAG	TTTGATTTTC	7320
ATTTACTAGA	AATGAAACTG	ATGAGAGATA	TCAGTAGACA	TTTGAGTCAG	GATATTATGG	7380
AAAATGATAA	AAAGAGCTCG	TGAGATTGGC	ATATCAGACT	ACTAAAGTAT	TGAGTTTGT	7440
AGGATTTTAG	CGACTAGTTA	GCTGGGAAAAG	GAAGATATTT	GTGACAAATA	ATAAACTGTA	7500
TTTCGTTGATA	GAATTTAGAA	ATAAAATATA	TGAAGAATTA	GAACTTTCCA	GAAGTGATTT	7560
AGCGATTTTA	CTATGTGCCA	TGCTTATCGC	CTCTATCGGA	TTAAATATGG	ATTCGACTCC	7620
CGTGATTATT	GGAGCCATGT	TAATCTCTCC	TTTGATGACA	CCTATTCTGG	GAGTGGGGCT	7680
CTCTCTAGCT	ATATTTGATT	TTAAATTGTT	AAGAAAATCT	TTTAAAATAT	TAGCTATTCA	7740

819

AATTCCTGCC AGTCTAATAG CTTCAACACT TTATTTTAT CTTTCTCCA TTTCGTATGC	7800
TAGTTCGGAG ATTGTTGCTA GAACCTCTCC GACTATTTGG GATGTTCTCA TTGCTTTTGT	7860
AGGAGGGATA GCAGGTATCA TTGGTGCTAG GAAAAAAGAG AC	7902

(2) INFORMATION FOR SEQ ID NO: 113:

- (i) SEQUENCE CHARACTERISTICS:
- (A) LENGTH: 18627 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 113:

GAAGTTGAAA TGGCCAGCTG ATGAGCAATA TCGGTCATAG AAATCTTCTC AATCAACTTT	60
TGCGCAATTT TTTGGTTGAT AATACGAGGA ATTTGGTGAT TTTTCTTGAC GATAGAAGTT	120
TCAGCGACCA TCATTTTGA ACAGTGATAG CACTTGAAAC GACGCTTCT AAGTAGAATT	180
CTAGTAGGCA TACCAGTTGT CTCAAGGTAA GGAATCTTAG ACGGTTTTG AAAGTCATAT	240
TTCTTCAATT GGTTCGCCA CTCAGGCCAA GATGGGGCGT CGTAGTCCAG TTTGGCGATG	300
ATTCCTTGT GTGTATCTT ATTGATGATG TCTAAATCT GGATATTAGG GTCTTTAATG	360
TCTAGTAATT TTGTGATAA ATGTAATTGT TCCATATGAA TCTTTCTAAT GAGTTGTTG	420
GTCGCTTTC ATTATAGGTC ATATGGGACT TTTTCTTCTAC AATAAAATAG GCTCCATAAT	480
ATCTATAAGG GATTTACCCA CTACAAATAT TATAGAGCCA AAAATCCTTT GTTTACTAAA	540
CAAGGGATTT TTCTTTTGT TCTGCTCCTT TTTTGATATA ATAGTTCTAT GTTAAATCA	600
GAAAAACAAT CACGTTATCA AATGTTAAAT GAAGAATTGT CCTTCCTATT GGAAGCGAA	660
ACCAATGTTT TGGCTAATCT TTCCAACGCC AGTGCTCTCA TAAAATCAG TTTTCCTAAT	720
ACCGTATTTG CAGGCTTTTA TTTGTTCGAT GGAAAGGAAT TGGTTTTAGG CCCCTTCCAA	780
GGAGGTGTTT CCTGCATCCG TATTGCACTA GGCAAGGGTG TTTGTGGTGA GGCAGCTCAC	840
TTTCAGGAAA CTGTTATTGT TGGAGATGTG ACGACCTATC TCAACTATAT TTCTTGATG	900
AGTCTAGCTA AAAGTGAAT TGTGGTGCCG ATGATGAAGA ATGGTCAGTT ACTTGAGTT	960
CTGGATCTGG ATTCTCAGA GATTGAGGAT TACGATGCTA TGGATCGAGA TTATTGGAA	1020
CAATTTGTCG CTATTTTGCT TGAAAAGACA GCATGGGACT TTACGATGTT TGAGGAAAA	1080
TCTTAATGTA TCAAGCACTT TATCGAAAAT ATAGAAGTCA AAATCTCTCC CAGTTAGTTG	1140
GTCAAGAAGT TGTGGCTAAG ACTCTTAAAC AAGCGGTGGA GCAAGAGAAA ATAAGTCACG	1200

CTTATCTTTT	TTCTGGTCCT	CGTGGAACGG	GAAAAACCAG	TGTTGCTAAA	ATCTTTGCCA	1260
AGGCTATGAA	CTGTCCCAAT	CAAGTGGGTG	GCGAACCTTG	CAATAACTGC	TATATTTGTC	1320
AAGCAGTGAC	GGACGGTAGT	TTAGAAGATG	TCATTGAAAT	GGATGCAGCT	TCTAATAATG	1380
GGGTAGATGA	AATTCGCGAA	ATTTCGTGATA	AATCTACCTA	TGCGCCTAGC	CTTGCTCGTT	1440
ATAAGGTTTA	TATCATAGAT	GAGGTTTACA	TGCTGTCTAC	AGGGGCTTTT	AATGCCCTCC	1500
TAAAGACGCT	GGAAGAACCA	ACACAGAATG	TAGTCTTTAT	TTTGCCACT	ACTGAATGTC	1560
ACAAGATTCC	TGCTACTATT	CTATCCCGTG	TGCAACGTTT	TGAGTTTAAA	TCAATTAAGA	1620
CACAGGATAT	TAAGGAACAT	ATTCACTATA	TCTTAGAAAA	AGAAAATATC	AGTTCTGAAC	1680
CAGAGCTGT	GGAAATCATT	GCCAGACGGG	CGGAAGGTGG	AATGCGGGAC	GCCTTGCTTA	1740
TTTTGGATCA	AGCCCTGAGT	TTGACACAGG	GAAATGAGCT	GACGACTGCT	ATCTCTGAAG	1800
AAATTACTGG	CACCATTAGC	CTATCAGCCT	TGGATGATTA	TGTGGCGGCC	TTGTCTCAAC	1860
AGGATGTTCC	CAAAGCTTTG	TCTTGCTTGA	ATCTTCTTTT	TGACAATGGT	AAGAGCATGA	1920
CTCGTTTTGT	GACCGATCTT	TTGCACTATT	TAAGAGACTT	GTTAATGTT	CAAACAGGGG	1980
GAGCAAATAC	TCATCATAGT	TCAGTCTTTG	TAGAAAATTT	GGCACTTCCT	CAAAAAATC	2040
TGTTTGAAAT	GATTTCGCTTA	GCAACAGTGA	GTTTAGCAGA	TATTAAGTCT	AGTTTGCAAC	2100
CCAAGATTTA	TGCTGAAATG	ATGACCGTCC	GTTTGGCGGA	AATCAAGTCC	GAACCAGCTC	2160
TATCAGGAGC	GGTTGAAAAT	GAAATGCTA	CGCTGAGACA	GGAAGTTGCC	CGTCTCAAAC	2220
AAGAGCTTTC	TAATGTAGGT	GCGGTTCCCTA	AACAAGTTGC	ACCAGCTCCT	AGTCGACCAG	2280
CTACGGGCAA	AACAGTCTAT	CGTGTGATC	GCAATAAAGT	GCAATCTATC	TTACAAGAGG	2340
CCGTCGAAAA	TCCTGATTTA	GCACGTCAAA	ATTTAATTCG	TTTGCAGAAT	GCCTGGGGAG	2400
AGGTAATTGA	AAGTCTAGGT	GGGCCGGACA	AGGCTCTGCT	AGTTGGTTCT	CAACCGGTTG	2460
CTGCCAATGA	ACACCATGCT	ATTCTTGCTT	TTGAGTCTAA	CTTCAATGCT	GGTCAAACCTA	2520
TGAAACGAGA	CAATCTCAAT	ACCATGTTTG	GTAATATCCT	CAGTCAGGCG	GCAGGTTTTT	2580
CACCTGAGAT	TTTAGCTATT	TCCATGGAGG	AATGGAAAGA	AGTTCGCGCA	GCCTTTTCAG	2640
CCAAAGCCAA	ATCTTCTCAA	ACTGAAAAAG	AAGTAGAAGA	AAGCCTGATT	CCAGAAGGAT	2700
TTGAATTTTT	GGCTGATAAA	GTGAAGGTAG	AGGAAGACTA	AAGAAAGATT	TCATGATACA	2760
ATAAGTTTAT	GAATAAACAA	CAATTATTATA	TTATGGCGCT	GTTTACAGCT	GCTGAGACCT	2820
ATTTTTTCAA	TGAAGCCTGG	ATGACTGGCC	GCTATATTAT	GGCAGCCTTT	TGGCAATTT	2880
TACTCTTTAG	AAATTTCCGA	GTCAGTTATG	TGATGGGCAA	AATCGTTGAT	GTCATCGATC	2940
AGCATTTTAA	TAGGAAAGAC	TAGCCCTCAG	CTTCCAGACA	AAATCAAAGC	CTTTTAGGCT	3000

TTTTTTTGTT	ATACTAGAAA	AGTATATTTA	TAGAATTTTT	GCTCTATTTT	TGGGGAAATC	3060
AGACGTTTTT	CTAGTAAGTA	CTGTAAAAGT	TTTGAAAAAG	AAAGGAACTA	TCATGTCACT	3120
ATTAGAGATC	AAAGATCTTC	ACGTTGAGAT	TGAAGGAAAA	GAAATTTTAA	AAGGGGTAA	3180
CCTGACCCTG	AAAACAGGAG	AAATTGCCGC	TATCATGGGA	CCAAATGGTA	CAGGTAATC	3240
GACTCTTTCT	GCCGCTATCA	TGGGAAATCC	AAACTATGAA	GTAACATAAG	GTGAAGTTTT	3300
GTTTGATGGC	GTAACATCC	TTGAGTTGGA	AGTGGATGAG	CGTGC CGTA	TGGGACTTTT	3360
CCTTGCTATG	CAATACCCAT	CAGAAATCCC	TGGAATTACC	AATGCTGAGT	TTCTTCGTGC	3420
CGCTATGAAT	GCGGGTAAAG	AAGATGATGA	GAAGATTTCA	GTTCGTGAGT	TTATTACTAA	3480
GCTAGATGAA	AAAATGGAAT	TGCTCAACAT	GAAAGAAGAA	ATGGCAGAGC	GTTACCTCAA	3540
CGAAGGCTTC	TCTGGTGGTG	AGAAAAACG	CAATGAAATT	CTTCAACTTT	TGATGTTGGA	3600
GCCAACATTT	GCTCTTTTGG	ACGAGATFGA	CTCAGGTCTT	GATATTGACG	CTCTTAAAGT	3660
TGTGTCTAAA	GGTGTCAAATG	CCATGCGTGG	TGAAGGTTTT	GGTGTCTATGA	TCATCACTCA	3720
CTACCAACGT	CTTTTGAACT	ATATCACACC	TGATGTGGTA	CACGTGATGA	TGGAAGGTCG	3780
TGTTGTCCTT	TCTGGTGGTC	CAGAATTGGC	TGCGCGTTTG	GAACGTGAAG	GATACGCAAA	3840
ATTAGCTGAA	GAACCTGGCT	ACGACTACAA	GGAAGAATTG	TAATTCCTTC	GTATCTTTTA	3900
GGAGAAGTAA	ATGACTAGAG	AAAATATTAA	ACTTTTTTCA	GAAATGCACG	CTGAACCAAG	3960
CTGGTTGGCT	GATCTCCGTC	AAAAAGCTTT	TGACAAGATT	GAGACTTTGG	AATTACCAGT	4020
TATTGAGTGT	GTCAAATTCC	ACCGTTGGAA	TCTGGGTGAT	GGAACGATTA	CAGAAAATGA	4080
GCCATCAGCA	AATGTTCCAG	ATTTACACAGC	TTTAGATCAT	CACTTGAAGT	TGGTGCAAGT	4140
AGGAAC TCAA	ACTGTTTTCG	AACAACTCC	AGTTGAGTTA	GCTGAACAGG	GTGTTGTCTT	4200
CACAGACTTT	CACTCAGCTT	TAGAAGAAAT	TCCAGAGCTG	ATCGAAGAAT	TCTTCATGTC	4260
ATCTGTTAAG	TATGATGATG	ACAAGTTGGC	GGCTTACCAC	ACAGCTTACT	TTAACAGTGG	4320
TGCTGTACTC	TATATCCAG	ATAACGTAGA	AATCACAGAG	CCAATTGAAG	GAATTTTCTA	4380
CCAAGATAGC	GATAGCAATG	TGCCGTTTAA	CAAGCATATT	ATGATTATCG	TTGGTAAAAA	4440
TTCTAAGATT	AGTTATCTGG	AGCGTTTAGA	GTCACGCGGT	GAAGGAAGTG	ACAAAGCAAC	4500
TGCCAATATC	ACAGTGAAG	TGATTGCACG	TTCTGGTGCG	CAAGTCAAGT	TTGCTGCTAT	4560
CGACCGTCTA	GGTGAACAG	TCACTGCCTA	CATTAGCCGT	CGTGGTAAAT	TAGGCAACGA	4620
TGCAAGTATT	GACTGGGCTA	TCGGTGTTCAT	GAACGAAGGA	AATGTCGTTG	CTGATTTTGA	4680
TAGTGACTTG	ATTGGTAATG	GTAGCCATGC	TGACCTCAAG	GTTGTAGCTC	TTTCAAGTGG	4740

822

TCGTCAGGTA	CAAGGGATTG	ATACTCGTGT	AACTAACTAT	GGCTGCAACT	CAATCGGAAA	4800
CATTCTACAA	CATGGGGTTA	TCCTTGAAAA	AGCAACTTTG	ACTTTCAATG	GTATCGGCCA	4860
CATCATCAAG	GGTGCTAAGG	GAGCAGATGC	GCAACAAGAG	AGCCGTGTTC	TCATGCTTTC	4920
AGACCAAGCG	CGTTCAGATG	CTAACCCAAT	TCTTTTGATT	GATGAAAATG	ACGTAAGTGC	4980
AGGCCATGCA	GCCTCTATTG	GTCAGGTAGA	TCCAGAAGAT	ATGTACTACC	TCATGAGTCG	5040
TGGCTTGAT	AAGGCAACTG	CAGAGCGTTT	GGTTGTTCGT	GGTTTCCTTG	GATCTGTAT	5100
CGTGGAGATT	CCAGTCAAGG	AAGTTCGTGA	TGAAATGATT	GCAACTATCG	AAGAGAAATT	5160
GTCAAAACGC	TAAGGGGCG	CCTATGTTAG	ATGTAGAAGC	GATTCGCAAG	GATTTTCCAA	5220
TTTTAGATCA	GATTGTCAAT	GATGAACCTC	TGTTCTATCT	GGACAATGCT	GCGACGACAC	5280
AAAAACCACT	AGTAGTTCTG	AAAGCTATTA	ACAGCTACTA	TGAGCAGGAC	AATGCCAATG	5340
TTCACCGTGG	TGTCCATACC	TAGCGGAAC	GAGCGACAGC	TTCTTATGAA	GCTGCTCGTG	5400
AAACCATTCTG	TAAGTTTATT	AATGCAGGCT	CTACAAAGGA	AGTTCTCTTT	ACCAGAGGAA	5460
CGACAACCAG	CCTTAAGTGG	GTGGCACGCT	TTGCTGAGGA	AATTCTCACT	GAGGGAGACC	5520
AGGCTCTGAT	TTCAGTAATG	GAACACCATT	CTAATATCAT	TCCATGGCAG	GAAGCTTGTC	5580
GAAAGACTGG	AGCAGAGCTT	GTCTATGTCT	ATCTTAAAGA	CGGTGCCTTG	GATATGGAGG	5640
ATTTGCGAGC	TAAATTGACT	GATAAGGTTA	AATTTGTFTC	CCTAGCTCAT	GCCTCCAATG	5700
TTCTTGGTGT	GGTCAATCCG	ATCAAGGAAA	TCACTCAATT	AGCCACCAA	GTGGGGCAA	5760
TTATGGTAGT	GGATGGTGCT	CAATCTACAC	CTCATATGAA	GATTGATGTC	CAGGACTTGG	5820
ATCTGGACTT	TTTCGCCTTT	TCGGGTCA	AGATGGCTGG	TCCGACTGGT	ATCGGTGTCC	5880
TTTACGGCAA	AGAAAAGTAT	CTTGAGCAAA	TGTCTCCAGT	AGAATTTGGC	GGCGAGATGA	5940
TTGATTTTGT	CTACGAGCAA	TTTGCTAGTT	GGAAGGAATT	GCCTTGAAA	TTGAGGCTG	6000
GAACGCCAAA	TATGGCAGGA	GCTATTGGAC	TTGCGACTGC	AGTTGATTAT	CTGGAAAAGA	6060
TTGGTATGGA	TGCCGTTGAA	GCTCATGAAC	AGGAATGAT	TGCGTACGTC	TATCCAAAAC	6120
TGCAGGCAAT	TGAGGGATTG	ACCATTTACG	GTTCTCAGGA	TTTGGCTCAA	CGTTCGGGTG	6180
TTATTGCCTT	TAACCTAGGT	GATCTCCATC	CTCACGATCT	TGCGACGGCT	CTGGATTATG	6240
AAGGAGTGGC	TGTTTCGTGCT	GGTCACCATT	GTGCGCAACC	CTTGCTTCAG	TATTTGGAAG	6300
TCCCAGCAAC	AGCTCGTGCA	AGTTTTTATA	TCTACAATAC	CAAGGCAGAT	TGCGACAAAC	6360
TAGTCGATGC	CCTACAAAAG	ACAAAGGAGT	TTTTCAATGG	CACTTTCTAA	ACTAGATAGC	6420
CTTTATATGG	CAGTGGTAGC	AGACCATTTCG	AAAAATCCAC	ATCACCAGG	GAAGTTAGAA	6480
GATGCTGAGC	AAATCAGTCT	CAACAATCCG	ACTTGTGGGG	ATGTCATCAA	CCTCTCTGTC	6540

AAGTTTGATG CAGAGGACCG TTTGGAAGAT ATTGCTTTTC TAAATTCAGG ATGCACGATT 6600
 TCAACTGCTT CTGCTAGTAT GATGACAGAT GCCGTTTTAG GAAAAACCAA ACAAGAAATT 6660
 TTAGAACTGG CACTATTTT TTCTGAAATG GTTCAAGGGC AAAAAGATGA GCGTCAAGAC 6720
 CAACTTGGAG ACGCGGCATT CTTGTCAGGT GTTGCCAAAT TCCCTCAAAG AATCAAGTGT 6780
 GCAACCCTAG CTTGGAATGC CCTTAAGAAA ACAATTGAAA ATCAAGAAAA ACAGTAAGAC 6840
 AAGTTTCTTT TGTCTTATGA ATTATTAGAA ATGAAGAAAG AAAGGATACT ATGGCTGAAG 6900
 AAAGAGTAGA ACCAAAACCA ATTGACCTTG GTGAATATAA ATTTGGTTTC CATGACGATG 6960
 TAGAGCCTGT CTTATCGACA GGAAAAGGAC TCAACGAAGG TGTTATTCGT GAATTATCTG 7020
 CTGCTAAGGG TGAGCCTGAG TGGATGTTGG AGTTCGGTTF GAAGTCTTAT GAAACCTTCA 7080
 AAAAAATGCC CATGCAAAC TGGGGAGCAG ACTTGTCAGA GATTGACTTT GATGACTTAA 7140
 TCTACTACCA AAAACCATCT GACAAAACCAG CCCGTTCTTG GGATGATGTA CCTGAAAAGA 7200
 TTAAAGAAAC CTTTGAACGT ATCGGGATTC CAGAAGCTGA ACGTGCTTAT TTAGCAGGGG 7260
 CTTCTGCCCA GTACGAGTCA GAAGTGGTTF ACCACAACAT GAAGGAAGAG TTCCAAAAAT 7320
 TAGGTATTAT CTTTACAGAT ACAGATTCCG CACTCAAGGA ATACCCAGAC TTATTTAAAC 7380
 AATACTTTGC GAAGTTGGTA CCGCCGACAG ATAACAAGTT GGCAGCCCTC AACTCAGCAG 7440
 TATGGTCGGG TGGAACTTT ATCTACGTGC CAAAAGGTGT CAAGGTAGAT ATTCCACTTC 7500
 AACTTATTT CCGTATCAAT AACGAAAATA TAGGTCAGTT CGAACGTACC TTGATTATCG 7560
 TTGATGAGGG AGCAAGCGTC TACTACGTAG AAGGATGTAC AGCACCAACA TATTCAAGCA 7620
 ATAGCTTACA CGCTGCCATT GTAGAAATTT TTGCTTTGGA CGGAGCTTAT ATGCGTTATA 7680
 CAACTATCCA AACTGGTCT GATAACGTCT ATAACCTGGT AACAAAGCGT GCTAAGGCTC 7740
 AAAAGGATGC CACTGTTGAG TGGATTGATG GAAACTTGGG TGCCAAAACG ACTATGAAAT 7800
 ATCCATCTGT TTACCTTGAT GGAGAAGGAG CGCGTGGTAC CATGCTCTCT ATCGCCTTTG 7860
 CTAATGCAGG GCAACACCAA GACACGGGTG CTAAGATGAT TCACAATGCT CCACATACCA 7920
 GCTCGTCTAT TGTGTCTAAA TCCATCGCTA AAGGTGGAGG AAAGGTTGAC TACCGTGGAC 7980
 AAGTCACCTT TAACAAGAAC TCTAAGAAAT CTGTTTCCCA CATTGAATGT GATACCATTA 8040
 TCATGGATGA CTTGTCAGCA TCAGATACTA TTCCATTTAA TGAAATTCAC AACTCGCAAG 8100
 TGGCTTTGGA ACACGAAGCC AAAGTATCTA AGATTCAGA AGAGCAATTG TATTATCTCA 8160
 TGAGCCGTGG ATTGTCAGAA TCTGAGGCAA CTGAAATGAT TGTGATGGGA TTTGTAGAAC 8220
 CCTTTACAAA AGAACTTCCA ATGGAATACG CAGTTGAGCT GAACCGCTTG ATTAGCTATG 8280

AAATGGAGGG	ATCAGTTGGA	TAAAATTTGA	TTTTATACTC	TTCGAAAATC	TCTTCAAACC	8340
ACGTCAGCAT	CGCCTTACCG	TATGTATGGT	TwCTGAtTCG	TCAGTTTCAT	CTACAACCTC	8400
AAAACAGTGT	TTTGAGCAAC	tGCGGCTAGC	TTCCTAGTTT	GTTCTTTGAT	TTTGAGTATT	8460
AGATTTACTC	AAAATCAAGG	ATTTTGAAGA	TGAACTTGTA	TCAAAAAATC	GCGGTTTAAA	8520
ATCGCGATTT	TTTATAATTT	CTCGTTAACA	AAGCGGACAA	ACTGATTCCA	CCAAACTTTT	8580
AAGAAGAAGG	CTTTTCAAT	TTTCTTGTCT	GCTACCATTT	CGAAACTAGG	GCGCTCTGTG	8640
GTGATGTAAC	CTTGACCAAT	CAAGTCCTTG	TCTTCATAAG	TCAAATGGCC	AACCACTGTT	8700
CCAGCTTCAA	GTGGTGCTGG	GATTGCTTTG	GAATCAGGTG	TGAATTGAAC	AGATTGGGAA	8760
GATTGATTCC	CAACACGTTT	GATTAGATAG	ATATCCTCTG	GAGCCACTGC	AGTTACTGTA	8820
TCTTCTTTTC	CATCTGTAC	AGGGGCTTTG	CTATCTTGAT	AGGCATCGCC	TTGTTGAACG	8880
ATTTTGCGAA	GTGTAAATGT	AGAAGAAATA	TAATCCATTA	GGGAAGATGT	AGCTGTAAAT	8940
CGAGCGTAAG	GATTATTGTC	TTGATGATCT	GCATTTAAAA	CAACTGTGAT	GACTCTCATG	9000
CCTTTTTCGA	CAGTAGTACC	AACAAAAGAC	TCTCCAGCCT	TATCTGTTGT	TCCTGTTTTT	9060
AGCCCATCAA	AACCACCACG	GTAAGCAGGC	ATACCTTCTA	ACATGTAGTT	GGTTGAAGTG	9120
ATTGTCATCC	CAGCAAAAGT	AGAAGAAGGT	TTTTTGGTGA	TTTCTAAGAC	TTGTGGGTAT	9180
TTTTTGATGA	GGTTGCGAGC	AACGATAGCG	ACATCATAAG	CACTAAGCTT	ATTTTCTCTA	9240
TCTTTTTTAG	AACCTGGGTA	AATGTTATCC	CCTAGAGTTT	CATTGTTAAG	ACCTGTGTA	9300
TTGACAACAG	TGGCATCCTG	AATTCCCCAT	TCCAAGAGTT	TTGCCCGCAT	CATATCGACG	9360
AAATCTTTTT	CTGAGCCAGC	AATTTTCTCA	GCTAGGGCAA	TAGCGGCGCT	GTTGGCACTA	9420
GATACCAGAG	TTGCTTCAAG	CAACTCTTCG	ACAGTATAAT	TACGGGCCTC	CATAGGAATA	9480
TTACTGGCTT	CAGAATTTGT	CGTCAATTGA	TAAGGATAAT	CAGAAATATC	TACAGGAGTG	9540
GAGAGGGTAA	TACTTCCGTT	TTCCAAAGCT	TCATAGACCA	GATAAACAGT	AATCAATTTT	9600
GTATGGAAG	CAATTCGAC	AGGTTGCGTT	GCATCCTTCT	CATAGAGAAT	TTTACCAGTA	9660
TTTGCCTCAA	CAGCAATCGC	ATGTTTAGCG	GCAATGGTAA	AATCTTGAGC	AACAGCAGTA	9720
GAAGCACCCC	CTAAAAGAGA	GACAGTTAAC	AAAGTTAAAA	ATATTTTTTT	CATAGTAGTC	9780
TTATTCTATC	ATAAAGAAAA	AAAATATTCT	TGCTTTAATA	ATTCATCTGT	TAAGCTTTTT	9840
GAAAATATGG	TAAAATAAAG	TAAGGGAGGT	AACTCATGTT	TCGTAGAAAT	AAATTATTTT	9900
TTTGACCAC	AGAAATTTTA	CTCTTAACCA	TCATCTTTTA	CCTATGGAGA	CAGATGGGGT	9960
CTTTGATTAA	CCCTTTTGTT	AGCGTGCTTA	ATACAATTAT	GATTCCATTT	TTATTAGGGG	10020
GCTTTTTTTA	TTATTTGACA	AACCCTATTG	TTACTTTCTT	AAATAAAGTC	TGTAAACTCA	10080

ATCGTTTGCT TGGTATTTTA ATTACCTTGT GFACTTTGGT CTGGGGAATG GTCATAGGTG 10140
 TTGTCTATCT CTTACCTATT TTGATTAATC AGTTATCTAG TTTGATTATA TCTAGTCAAA 10200
 CTATTTATAG TCGAGTACAA GACTTAATCA TAGACTTATC TAATTATCCT GCGCTCCAGA 10260
 ATTTGGATGT AGAAGCTACA ATTCAGCAGT TAAACTTATC CTATGTTGAT ATTCTTCAAA 10320
 ATATCCTAAA TAGCGTATCA AATAGTGTGG GGAGCGTCTT GTCAGCTCTT ATCAGTACTG 10380
 TTTTGATTTT GATTATGACT CCAGTTTTTT TGTTTTATTT CTTATTAGAT GGACATAAAT 10440
 TCTTGCCCAT GCTTGAAGA ACGATTCTAA AGAGGGATCG CTGTCATATT GCAGGCTTAT 10500
 TAAAGAATTT AAATGCGACG ATTGCTCGCT ATATTAGTGG AGTTTCGATT GACGCAATCA 10560
 TTATAGGTTG TTTGGCTTAT ATTGCTATA GTATTATTGG TTTAAAATAT GCTTTAGTTT 10620
 TTGCCATTTT TTCTGGTGTA GCCAATTTAA TTCCTTATGT GGGGCCAAGT ATTGGTTTGA 10680
 TTCCTATGAT CATCGCAAAT ATATTCCTG ATCCCCATAG ACTGCTGATT GCAGTGATTT 10740
 ATATGCTTGT TGTTTCAGCAG GTAGATGGCA ATATCTTATA TCCTCGAATC GTAGGAAGTG 10800
 TTATGAAGGT TCATCCAATC ACGATTTTAG TTTTACTTTT GTTGTCGAAGC AATATCTATG 10860
 GTGTAGTTGG AATGATTGTC GCAGTGCCAA CCTATTCTAT CTGAAAGAA ATTTCTAAGT 10920
 TCTTATCCCA TTTGTATGAA AATCATAAAA TAATGAAAGA ACGAGAAAGA GAATTAGCTA 10980
 AGTAAAAGTC AGGAGAACCC TGATTTTTCT TTAAGGAAAG TGGCCTTTAG ATTAGAAGAC 11040
 TGAAAATAAG TTAAAGTCTT AACTAATTT TCACAGCTAA GAATAGTAGA AGTTAATCTG 11100
 ATAAAAATCG AAAAAACCAG TGAATTCTG TGTCAGGGTA AGTTCCACTG GTTTTCATAG 11160
 TCTATTAAAG TTCGAATGAA ACCTATTTAT AGTAGATTGA AACTAGAATA GTACACCTCT 11220
 AATTCTAAAA CATTGTTAGA AATCGATTG ACTGTCCTGA TCTATTCGTT CTATTCCTAT 11280
 TTCATTTTAC TATATTTTGG TGCAATAAGT GAAAAGTAGT CCGAATAATA TAAGGATTGA 11340
 TTTTATAGTT TTTAACTCA AATGAATTGA AATAAAGAGA GTACGAAAAT TCTCATCTGA 11400
 AAGTATTTTA GAATAATCT CTTCGTGAAT TTCTTCAAAA CAGATAGCTT CATCTTAGGT 11460
 ATGTGATTTT TTTTGCATT TTTGAGTTAG ATAAGGTATA ATGATTTTAT TGTCTTTTGG 11520
 GGTTCGTTACG GATTCGACAG GCATTATGAG GCATATTTT CGACTCGTGT GGCGACGTAA 11580
 ACGCTCAGTT AAATATAACT GCAAAAAATA ACACTTCTTA CGCTCTAGCT GCCTAAAAAC 11640
 CAGCAGGCGT GACCCGATTT GGATTGCTCG TGTTCAATGA CAGGTCTTAT TATTAGCGAG 11700
 ATACGATTAA GCCTTGCTA GCGGTTTGGT AAGAGATTGA TAGACTCGCA GTTCTAGAC 11760
 TTGAGTTATG TGTCGAGGGG CTGTTAAAAT AATACATAAC CTATGGTTGT AGACAAATAT 11820

826

GTTGGCAGGT	GTTTGGACGT	GGGTTGCGACT	CCCACCGGCT	CCATTATTCC	TTTGCATTCT	11880
TTTGCATTCC	TTGGTAAAAC	GTTGTAAAT	CAACGTTTTT	TATTTTTATC	TTTGGTATTC	11940
CTTTGCATTC	TTTTGCTAAA	AAGGGAGTCA	CAAACAGACC	CTATTTTAAA	AAAGGATAGA	12000
AAAAAGGATA	CAACATTTGT	CGCATCCTAA	AAATAATCTT	TTTTTCGACGG	AAGACATGGG	12060
ATTCGAACCC	ACGCACGCTA	TTACACGCCT	ACCGCGTTTC	CAACACGGCC	TCTTAAGCCT	12120
CTTGAGTAAT	CTTCCAATAC	TTACTCAAAT	AGTCTACCAT	AAAGGCTCTT	ATCTTGCAAT	12180
AAAAATCTTA	GAAATAAGAA	AAATGATAGA	TTTTGAAAGA	AAATGATAAA	AAATGCTTGA	12240
CTTCGAAAGA	AAGTATGATA	GAATGAATAG	TGTAAACGAT	AACAGGAGGT	GATTCAGTGT	12300
TAAAAACAGA	ACGTAAACAA	CTAATTTTAG	AGGAGTTAAA	TCAACATCAT	GTAGTTTCTC	12360
TAGAAAAATT	AGTTAGTTTG	CTAGAAACGT	CAGAATCAAC	GGTTCGAAGA	GACTTGGATG	12420
AGTTGGAAGC	GGAAAACAAG	CTTCGTCGTG	TGCATGGTGG	AGCAGAACTC	CCCTACTCCT	12480
TACAGGAAGA	AGAAACCATT	CAAGAAAAAT	CTGTCAAAAA	CCTTCAAGAA	AAGAAATGTC	12540
TGGCTCAGAA	AGCAGCCTCT	CTCATTAAG	AAAAAGATGT	CATCTTTATC	GATGCTGGAA	12600
CAACAACCTGC	TTTTTTGATT	CATGAATTGG	TCAATAAGAA	TGTTACAGTT	GTGACCAACT	12660
CCATTCACCA	TGCCGCTCAG	TTGGTTGAAA	AGCAGAWTCC	AACTGTTCATG	GTTGGAGGAA	12720
ACGTCAAGAC	GGCGACAGAT	GCTAGTATCG	GGGGCGTTGC	TCTTAACCAG	ATTAACCAAT	12780
TGCACTTTGA	CCGTGCCTTT	ATCGGAATAA	ATGGTGTGTA	CGATGGCTAT	TATACGACTC	12840
CTGATATGGA	GGAGGGAGCT	GTGAAAAGAG	CTATTTTGGG	GAATGCCAAG	CAGACCTACG	12900
TCTTGGTGGG	TTCGTCAAAA	ATTGGACAAA	CTTGCTTTGC	CAAGGTAGCC	CCACTCAAAC	12960
GCGCTATCGT	TATCACTAGT	CAAGGGCATG	AGCTCTTGCA	GGTTATTAAG	GAGAAAACGG	13020
AGGTAATAGA	AGTATGATTT	ATACAGTCAC	ACTCAATCCA	TCCATTGACT	ATATCGTTTCG	13080
TTTGGACCAA	GTCAAAGTTG	GTAGTGTCAA	TCGTATGGAC	AGTGATGATA	AGTTTGCTGG	13140
TGGGAAAGGA	ATCAATGTCA	GCCGTGTCTT	GAAACGTTTG	AATATACCAA	ATACAGCGAC	13200
GGGATTTATC	GGTGGCTTTA	CTGGTAAATT	TATCACAGAT	ACTTTAGCAG	AGGAAGAAAT	13260
CGAGACACGT	TTTGTCCAGG	TGGCAGAAGA	TACTCGTATC	AATGTTAAAA	TCAAAGCAGA	13320
CCAAGAAACA	GAAATCAACG	GAACGGGTCC	AACTGTTGAA	TCGGTTCAGC	TAGAAGAATT	13380
GAAAGCTATT	TTATCTAGTC	TGACAGCAGA	AGATACAGTT	GTCTTTGCAG	GTTCAAGTGC	13440
TAAAAATCTA	GGCAATGTTA	TCTATAAGGA	TTTGATTTCC	TTGACGCGCC	AGACTGGTGC	13500
GCAAGTGGTC	TGTGACTTTG	AAGGACAGAC	CTTAATTGAT	AGTTTGGACT	ACCAGCCTCT	13560
TCTTGTAATA	CCAAACAATC	ATGAACCTGG	AGCGATTTTT	GGGGTTAAAC	TCGAAAGTTT	13620

AGATGAAATT GAGAAATACG CTCGTGAGTT ACTGGCTAAG GGTGCTCAA ATGTTATTAT	13680
CTCTATGGCT GGTGATGGTG CCCTTCTTGT CACATCTGAG GGAGCTTACT TCGCTAAACC	13740
AATCAAAGGA ACAGTCAAAA ATTCAGTTGG AGTGGTGAT TCTATGGTTG CTGGATTAC	13800
AGGTGAATTT GTCAAATCAA AAGACGTAGT AGAAGCCTTC AAATGGGGAG TGGCTTGCGG	13860
AACGGCAACT ACCTTCTCAG ATGACTTGGC AACGGCGGAA TTTATTAAAG AAACATATGG	13920
AAAAGTTGAG GTAGAAAAAC GATGAAAATT CAAGACCTAT TGAGAAAAGA TGTCATGTTG	13980
CTAGATTTGC AGGCAACTGA AAAACAGCT GTCATCGACG AGATGATTAA AAATTTGACA	14040
GACCACGGTT ATGTAACAGA TTTTGAAACA TTTAAAGAAG GAATTTTGGC GCGTGAAGCT	14100
TTGACTTCTA CTGGTTTGGG TGATGGAATC GCAATGCCTC ACAGCAAAAA CGCTGCTGTC	14160
AAAGAAGCGA CAGTTCATT TGCTAAGTCA AATAAGGGTG TTGACTACGA GAGCTTGGAT	14220
GGACAAGCAA CTGACCTCTT CTCATGATT GCAGCTCCAG AAGGTGCCAA TGATACTCAC	14280
TTGGCAGCCT TGGCAGAATT GTCTCAATAC TTGATGAAAG ACGGTTTTCG AGACAAACTT	14340
CGTCAAGCAA CATCTGCAGA CCAAGTTATC GAACTTTTTC ACCAAGCTTC AGAAAAAACT	14400
GAGGAACCTG TTCAAGCACC TGCTAATGAC TCTGGTGACT TTATCGTAGC TGTTACAGCT	14460
TGTACAACAG GTATTGCCA CACTTACATG GCCCAAGAAG CCCTTCAAAA AGTAGCTGCT	14520
GAAATGGGGG TTGGTATCAA GGTGCAAAAC AACGGTGCTA GCGGTGTTGG AAATCAACTA	14580
ACTGCAGAAG ATATCCGTAA GGCTAAAGCT ATTATCATTG CAGCAGACAA GGCCGTGAA	14640
ATGGATCGAT TTGATGAAA ACCATTGATC AATCGTCCAG TTGCTGACGG TATCCGTAAG	14700
ACAGAAGAGC TAATTAACCT GGCTCTTTCA GGAGATACTG AAGTCTACCG TGCCGCTAAT	14760
GGTGCCAAAG CTGCAACAGC CTCTAACGAA AAACAAAGCC TTGGTGGTGC CTTGTACAAA	14820
CACTTGATGA GTGGTGTATC TCAAATGTTA CCATTCGTTA TCGGTGGTGG TATCATGATT	14880
GCCCTTGCCCT TCTTGATTGA CGGTGCCTTG GGTGTTCCTA ATGAAAACCT TGGCAATCTT	14940
GGTTCCTFACC ATGAGTTAGC TTCTATGTTT ATGAAAATTG GTGGAGCTGC CTTTGGTTTG	15000
ATGCTTCCAG TCTTTGCGGG TTATGTTGCC TACTCTATTG CTGAAAAACC GGGTTTGTA	15060
GCAGGTTTCG TGGCTGGTGC TATTGCCAAA GAAGGTTTTC CCTTTGGTAA AATTCCTTAT	15120
GCCGCAGGTG GTGAAGCAAC TTCAACTCTT GCAGGTGTCT CATCTGGTTT CTFAGGTGCC	15180
CTTGTGGTG GATTTATCGC AGGTGCCTTG GTTCTTGCCA TCAAGAAATA CGTTAAAGTT	15240
CCTCGTTCAC TCGAAGGTGC TAAATCAATC CTTCTATTGC CACTTCTTGG AACAACTTG	15300
ACAGGATTTG TTATGCTAGC TGTGAATATC CCAATGGCTG CAATCAACAC TGCTATGAAT	15360

GACTTCCTAG	GCGGTCTTGG	AGGAGGTTCA	GCTGTCCCTC	TTGGTATCGT	CCTTGGTGGG	15420
ATGATGGCTG	TTGACATGGG	TGGACCAGTT	AATAAAGCAG	CTTATGTCTT	TGGTACAGGT	15480
ACGCTTGCAG	CAACTGT TTC	TTCAGGTGGT	TCTGTAGCCA	TGGCAGCAGT	TATGGCTGGA	15540
GGAATGGTGC	CACCACTTGC	AATCTTTGTC	GCAACTCTTC	TTTTCAAAGA	TAAATTTACT	15600
AAGGAAGAAC	GTAACCTCTGG	TTTGACAAAC	ATCATCATGG	GCTTGTCAAT	TATCACTGAG	15660
GGAGCGATTC	CATTTGGTGC	CGCTGACCCA	GCTCGTGCGA	TTCCAAGCTT	CATCCTTGGT	15720
TCAGCAGTAG	CAGGTGGACT	CGTTGGTCTT	ACTGGTATCA	AACTCATGGC	GCCACACGGA	15780
GGAATCTTCG	TTATCGCCCT	TACTTCAAAT	GCTCTCCTTT	ACCTCGTTTC	TGTCTTGGTA	15840
GGAGCAATCG	TAAGTGGTGT	GGTTTATGGT	TACCTACGCA	AACCACAAGC	ATAAAAAATA	15900
GAAAAATGAA	AAGATTGGAC	CGTTTGGTGC	AGTCTTTTTC	TCTTCCCGAA	ATGCCTGTGA	15960
AATATGGTAT	AATAGAAGAA	TGGCAAACAA	GAATACAAGT	ACAACAAGAC	GGAGACCCTC	16020
TAAAGCAGAA	CTGAAAAGAA	AAGAAGCGAT	TCAACGAATG	TTGATTTTCG	TAGGAATTGC	16080
GATTTTATTG	ATTTTCGCAG	CCTTCAAATT	AGGGGCTGCA	GGTATAACCC	TTTATAATTT	16140
AATTCGCTTG	CTAGTGGGTA	GCCTAGCTTA	TCTGGCGATA	TTCCGCCTAT	TAATCTATCT	16200
CTTCTTTTTC	AAGTGGATAC	GAAAACAGGA	AGGACTCTTA	TCTGGCTTTT	TCACCATATT	16260
TGCTGGCTTA	CTCTTGATTT	TTGAGGCCTA	CTTGGTTTGG	AAATATGGTT	TGGACAAGTC	16320
CGTTCTAAAA	GGGACCATGG	CTCAGGTTGT	GACAGATCTG	ACTGGTTTTTC	GAACGACTAG	16380
CTTTGCTGGA	GGGGGCTTGA	TCGGGGTCGC	TCTTTATATT	CCAACAGCCT	TTCTCTTTTC	16440
AAATATCGGA	ACTTACTTTA	TTGGTTCTAT	CTTGATTTTA	GTGGGTTCTC	TCCTAGTCAG	16500
CCCTTGGTCT	GTTTACGATA	TTGCTGAATT	TTTCAGTAGA	GGCTTTGCCA	AATGGTGGGA	16560
AGGGCACGAG	CTCGAAAAG	AGGAACGCTT	TGTCAAACAA	GAAGAAAAAG	CTCGCCAAAA	16620
GGCTGAGAAA	GAGGCTAGAT	TAGAACAAGA	AGAGACTGAA	AAAGCCTTAC	TCGATTTGCC	16680
TCCTGTTGAT	ATGGAAACGG	GTGAAATTCT	GACAGAGGAA	GCTGTTCAAA	ATCTTCCACC	16740
TATTCCAGAA	GAAAAGTGGG	TGGAACCAGA	AATCATCCTG	CCTCAAGCTG	AACTTAAATT	16800
CCCTGAACAG	GAAGATGACT	CAGATGACGA	AGATGTTTCT	GTCGATTTTT	CAGCCAAAGA	16860
AGCCCTTGAA	TACAAACTTC	CAAGCTTACA	ACTCTTTGCA	CCAGATAAAC	CAAAAGATCA	16920
GTCTAAAGAG	AAGAAAATTG	TCAGAGAAAA	TATCAAAATC	TTAGAAGCAA	CCTTTGCTAG	16980
CTTTGGTATT	AAGGTAACAG	TTGAACGGGC	GGAATTTGGG	CCATCAGTGA	CCAAGTATGA	17040
AGTCAAGCCG	GCTGTTGGTG	TAAGGGTCAA	CCGCATTTCC	AATCTATCAG	ATGACCTCGC	17100
TCTAGCCTTG	GCTGCCAAAG	ATGTCCGGAT	TGAAGCACCA	ATCCCTGGGA	AATCCCTAAT	17160

CGGAATTGAA GTGCCCAACT CCGATATTGC CACTGTATCT TTCCGAGAAC TATGGGAACA	17220
ATCGCAAACG AAAGCAGAAA ATTTCTTGGA AATTCCTTTA GGAAGGCTG TTAATGGAAC	17280
CGCAAGAGCT TTTGACCTTT CTAAAATGCC CCACTTGCTA GTTGACAGTT CAACGGGTTT	17340
AGGGAAGTCA GTAGCAGTTA ACGGCATTAT TGCTAGCATT CTCATGAAGG CGAGACCAGA	17400
TCAAGTTAAA TTTATGATGG TCGATCCCAA GATGGTTGAG TTATCTGTTT ACAATGATAT	17460
TCCCCACCTC TTGATTCCAG TCGTGACCAA TCCACGCAA GCCAGCAAGG CTCTGCAAAA	17520
GGTTGTGGAT GAAATGGAAA ACCGTTATGA ACTCTTTGCC AAGGTGGGAG TTCGGAATAT	17580
TGCAGGTTTT AATGCCAAGG TAGAAGAGTT CAATTCCTCAG TCTGAGTACA AGCAAATTC	17640
GCTACCATTC ATTGTCGTGA TTGTGGATGA GTTGGCTGAC CTCATGATGG TGGCCAGCAA	17700
GGAAGTGAA GATGCTATCA TCCGTCTTGG GCAGAAGGCG CGTGCTGCAG GTATCCACAT	17760
GATPCTTGCA ACTCAGCGTC CATCTGTTGA TGTCATCTCT GGTGTGATTA AGCCAATGT	17820
TCCATCTCGT GTAGCATTTC CGGTTTCATC AGGAACAGAC TCCCGTACGA TTTTGGATGA	17880
AAATGGAGCA GAAAACTTC TTGGTCGAGG AGACATGCTC TTTAAACCGA TTGATGAAAA	17940
TCATCCAGTT CGTCTCCAAG GCTCCTTTAT CTCGGATGAC GATGTTGAGC GCATTGTGAA	18000
CTTCATCAAG ACTCAGGCAG ATGCAGACTA CGATGAGAGT TTTGATCCAG GTGAGGTTTC	18060
TGAAAATGAA GGAGAATTTT CGGATGGAGA TGCTGGTGGT GATCCGCTTT TTGAAGAAGC	18120
TAAGTCTTTG GTTATCGAAA CACAGAAAGC CAGTGCCTCT ATGATTCAGC GTCGTTTATC	18180
AGTTGGATTT AACCGTGCGA CCCGTCTCAT GGAAGAAGT GAGATAGCAG GTGTCATCGG	18240
TCCAGCTGAA GGTACCAAAC CTCGAAAAGT GTTACAACAA TAAAAAATA GCTTCTTTCC	18300
AAGTTTGAG GGAAGCTATT TTAGTGGCTA TTGATTGCTT TTATTTTCTG AAGTTGGCGC	18360
ATPGGACTGT TTTTCGTTTT CAGTAGCAGG TTTACTTGAA GCAGGAGTAG AAGAGTCCTG	18420
AGTTGCTGTT TTCTGATCTT CTTTTTCTC TTCCTTGACG CTAGATTTTG GTGTTCCCTC	18480
TGCTGTGTT TTTTCTTGAC TAGTGTTAGT CTCPTTAGTT GACTGGTGT TTTCCTTAGG	18540
GGATTCCTTT TGATTTCTT TGACAATGTT TGTCGTCTGG CTGTCCGTAG GTTCTTTTTT	18600
AATATTTTGG TTATTATCCA AGGCGTT	18627

(2) INFORMATION FOR SEQ ID NO: 114:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 2560 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 114:

TAAAATACGT TACCTTGCTT CTGCACG TTC AGCAGGTAAG TCATTGAAAT TTAAAGATCA	60
AGATATTACA ATTGAAGAAA CGACTGAAAC AGCTTTTGAA GGAGTTGATA TTGCTCTCTT	120
TTCAGCAGGT AGTTCTACAT CAGCTAAGTA TGCACCATAC GCAGTAAAAG CTGGCGTGGT	180
AGTAGTAGAT AATACATCTT ATTTCCGTCA AAATCCAGAT GTTCCCTTGG TTGTTCCAGA	240
GGTCAATGCT CATGCACTTG ATGCTCACAA CGGAATCATT GCCTGCCCTA ATTGTTCAAC	300
AATTCAAATG ATGGTGGCTC TTGAGCCGGT TCGCCAAAAA TGGGGCTTGG ACCGTATCAT	360
TGTTTCAACT TATCAAGCCG TTTCAGGTGC TGGTATGGGA GCAATTCCTG AGACACAACG	420
TGAACTTCGT GAAGTCTTGA ATGATGGTGT GAAACCACGT GATTTGCATG CGGAAATCTT	480
GCCTTCAGGT GGTGACAAGA AACATTATCC TATCGCCTTT AACGCTCTTC CACAAATTGA	540
TGTTTCACT GATAATGATT ACACGTACGA AGAGATGAAG ATGACCAAGG AAAC TAAGAA	600
AATTATGGAA GATGATAGCA TTGCAGTATC TGCAACATGT GTGCGTATTC CAGTCTTGTC	660
AGCTCACTCT GAGTCTGTTT ATATCGAAAC AAAAGAAGTG GCTCCAATCG AAGAAGTAAA	720
AGCAGCTATC GCAGCCTTCC CAGGTGCTGT TCTTGAAGAT GATGTAGCTC ATCAAATCTA	780
TCCTCAAGCT ATCAATGCAG TTGGTTCGCG TGATACCTTT GTTGGTCGTA TCCGTAAAGA	840
CTTGATGCA GAAAAAGGAA TTCACATGTG GGTGTTTCA GATAACCTTC TCAAAGGTGC	900
TGCTTGGAAC TCAGTTCAGA TTGCTGAAAC TCTTCATGAA CGTGGATTGG TTCGTCCAAC	960
AGCCGAATTG AAATTTGAAT TAAAATAGTC ATATCGTTTA GGAGTTCAGA TGAATCCTT	1020
CTTTGAAATA GAGAGGTGTT TTCGTGTCTT ATCAAGATTT AAAAAAATGT AAAATCATT	1080
CAGCCTTTAT TACCCCTTC CATGAGGATG GTTCCATTAA CTTTGATGCT ATTCCAGCCT	1140
TGATFGAGCA TTTATFGGCC CATCATACGG ATGGAATCT TCTCGCAGGA ACGACTGCTG	1200
AGAGTCCAAC TTTGACCCAC GATGAGGAGT TGGAGTTGTT TGCGGCTGTA CAAAAGGTTG	1260
TCAATGGACG CGTTCCTTTG ATTGCGGGTG TAGGTACTAA TGATACGCGT GACTCTATTG	1320
AGTTTGTCAA AGAAGTAGCG GAATTTGGTG GTTTCGCAGC TGGGCTTGCT ATTGTTCTT	1380
ACTACAACAA ACCTTCCTCAA GAAGGATGT ATCAGCACTT TAAGACTATT GCAGATGCTT	1440
CTGACCTACC AATTATTATC TATAACATTC CAGGGCGTGT AGTTGTGCGA TTGACTCCAG	1500
AAACCATGCT TCGCTTGCT GACCATCCAA ATATTATCGG TGTCAAAAGAA TGTACTAGCT	1560
TGGCTAATAT GGCTTACTTG ATTGAGCACA AGCCTGAAGA GTTCTTGATT TATACAGGTG	1620
AGGATGGAGA TGCTTTCCAT GCCATGAACC TTGGGGCGGA TGGGGTTATT TCTGTTGCCT	1680

831

CTCATACAAA TGGGGATGAA ATGCACGAGA TGTTTACTGC GATTGCAGAA AGCGATATGA	1740
AGAAAGCCGC AGCAATTCAG CGTAAATTC AATGCTCTC TTCTTTTATC	1800
CAAGTCCTGC TCCAGTTAAG GCAATTCTTA ACTATATGGG ATTTGAAGCT GGACCCACTC	1860
GTCTACCTCT TGTTCAGCA CCAGAAGAAG ATGCCAAACG CATTATCAAG GTTGTCGTAG	1920
ATGGCGACTA CGAAGCAACT AAGGCAACTG TAACAGGGGT CTTAAGACCA GATTACTAAT	1980
AAAGACAATA AATCCGGCT CTTTGTCAAC TG TAGTGGT TGAAGTCAGC TAAGCTCGAG	2040
AAAGGACAAA TTTTGTCTT TCTTTTTGA TATTCAGAGC GATAAAAATC CGTTTTTTGA	2100
AGTTTTCAAAA GTTCCGAAAA CCAAAGGCAT TGCCTTGAT AAGTTTGATG AGATTATTGG	2160
TCGCTTCCAA TTTGGCGTTT GAATAGGGTA GTTGAAGGGT GTTGACGATT TTCTTTTTGT	2220
CCTTTAGAAA GTTTTTAAAG ACAGTCTGAA AATAGGATG AACCTGCTTC AGATTGTCTT	2280
CAATGAGTCC GAAAAATTC TCCGGTCTT TATTTCTGAAA GTGAAACAGC AAGAGTTGAT	2340
AGAGCTGATA GTGATGTTT AAGTTTTGTG AATAGCTCAA AAGCTTGTTT AAAATCTCTT	2400
TATTGGTTAA GTGCATACGA AAAGTAGGAC GATAAAATCG CTTATCACTC AGTTTACGGC	2460
TATCCTGTTG AATGAGTTT CAGTAGCGCT TGATAGCCTT GTATTCCGGGA TTTTCGATGA	2520
AACTGATTCA TGATTTGGAC ACGCACACGA CTCATAGCAC	2560

(2) INFORMATION FOR SEQ ID NO: 115:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 11303 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 115:

TATTGGATTT CCCTTGCAAT CAGTTTATGG GACAAGCACC CGGCAGCGCA GAGGAAATCA	60
ACGCCTTCTG TAGCCTACAT TTCAAACCA CCTTCCCACG TTTTGCCAAG ATTAAGGTCA	120
ACGGTAAGGA AGCAGACCCT CTCTATGTCT GGTACAAGA CCAGAAATCC GGCCACTAG	180
GAAAACGAGT CGAATGGAAT TTCGCTAAGT TTCTCATCGG TCGAGATGGG CAAGTCTTTG	240
AACGCTTTTC TTCAAAAACA GACCCAAAAC AAATTGAAGA GCGATACAA ACTCTACTAT	300
AATTCACAAT CTCACTATGA TTAGGTTTCC TTTAACCTGA TGAATAGTGA GATTTTTTGA	360
TGGGCTTTGA CTTAAATAGA AAAACACCCC ATGATATGAA ACATGAAGTG TTGTAAAGTC	420
TATGTTGTAG GTGCTTATTT CACAATTCA ATGTGACCAG TGATAACGAA TACCATACAG	480

AATCTTCATA	TACACTAAAC	AAATGACTTT	CTAATTATTT	CAATTAGTTT	TGGCTAGTAA	540
ATATCATTTC	CAACAAACGC	CCTCTCAATT	CCTTATCCTG	ATGATGCAAG	ATATTCATTA	600
AGTCATGAGA	GTTTTTCGCA	TTGATGAATT	GATTTAACAA	TCTATCTTTT	AATTCATATG	660
GAAGAGAAGC	TGCTTTTAGT	AGTCTAAAAA	CTTCGTCATT	TAAAGATGTC	CTTTTATTAT	720
CTTTCCATTC	AAATTTAGCT	GTATCATTCT	TATTTGGCAA	TTCAATTATA	GACACATTCG	780
TTCCTTTAAA	ATGAATTCTA	TGTTTTCTAT	TGCTTGGAAC	GATACTAGAA	TCTCCTTGTA	840
ATGCTAACTC	TACCATTCCC	ATTTCCCAAT	CGATTGATAA	TCTTGTTTTA	TATCTTTGAC	900
CATTTTGATC	TTCAAGCATT	TCAAAAGAAT	GTTGTTTTCC	TGGGAATACA	TACCAATCTA	960
CAACTTCAGG	TAAATCAACA	CCCATACCTA	TCTCAGAACC	AACCAAGGGA	ATGATTGCAC	1020
CACTTTTTGC	AAACACAGGC	GTAGTCGAGA	TGTCCTATA	AACACTTAAC	TTCACACCAC	1080
CTGTGTATTT	TTTCTCTGAA	AAGAAGTCAT	ACCATTCCACC	TTCAGGGAAC	CATACATCTA	1140
CTTTTGCAGA	TTGGAATGTC	AAATCCATCT	TTTCTACAAT	GGGAGCCACC	ATCAGTTCTG	1200
TTCCAAAAAA	GTATTGGTTT	GGAACATTAT	AGCTCTCATC	ATTCTCTGGA	TAGAAATAAT	1260
AGATTGGACT	GATTAATGGG	GCACCTTCCT	CATGTGTCTG	TACATTCATG	GTATATAGAT	1320
AGGGAATCAT	CTGATGTCTC	AAACGAAGGT	ATTTCTTCAT	AATCTTAGAT	GTTGTTTCTG	1380
AAAAAAACCA	AGGTTCCTTA	CTATTAAAAG	GACTTCTAGA	ACTATGTAAT	CGAGTAATCG	1440
GACTAAAAAC	ACCAAACTGT	AGCCATCTAG	TTTGTAGCTC	TTCGTCATAA	TCCCCAACAA	1500
TATGTCCACC	GATATCATGA	CTCCACCAAC	TATAACCGAT	ATTAGATGCT	GTCGCTGTAA	1560
AATAGGGTTG	AAATCTTAAG	GAATTCCAAC	TAATAATAGT	ATCCCCTGAA	AAACCAACAG	1620
GGTAGCGGTG	ACTACCAGGA	CCTGCATATC	TTGATAAAAT	CAAACCACCT	TCTGCATTTT	1680
TACAACATATC	CTGATAGTGA	TAATGGTTTA	AAAGCCAAAG	TGGATCTAGC	ATACCTTGTG	1740
TCCCTTGTGTG	CCAGTCAATC	CACCAAAAAT	CTACTCCCTG	CTTTTCTAGT	TCATAATGAA	1800
CATCTTTAAA	GTAGGCTTCC	CTAAAAGAGG	GATTAAAAAA	ATCAAAAATA	GCAGGTTCTT	1860
CTAGTTCTAC	ATTTAACCCC	AACCGTTTTG	CGATTTGAGG	ATAAGCTTCT	TCATAAGCCC	1920
GTATCCCATC	AGCAGGATGG	ACATTTAAGG	AGAGTTTTAG	CTTCTATCA	TGAAGTTGTT	1980
GCAATAACTG	TTCTGGATTT	GGTATTAAGT	TTCTATTCCA	ACTATATCCT	GTCCAGCCAC	2040
TTCCAAAGCG	AGCTGGAATG	TCAGTTATAT	GCCAATCCAT	ATCTAACACA	CCGATAGATA	2100
ATGGAATTTT	CTCTGTTTCA	AATCTGTCTA	TTAAATCCAA	GTATTCATCC	GACGTATAAG	2160
GCCAATATCT	ACTCCACCAA	TTGCCTAAAG	CATATCTTGG	CAACAAGGGT	GTTGAACCAG	2220
TCAAATGGTA	AAAATCTCTG	ATTGCTCCTC	TATAATCATG	CCCATAGGCA	AAGAAATACA	2280

833

GGTCAATTTG	ATTTTCTCTC	TCAATATAAC	CAGATTGTTC	ATCCCAAATA	AATCCTTGAG	2340
AATCATCCAA	TAAGGCTATA	CCATTTGGC	TAATAATTCC	ATCTTCTAAC	GAGATTGCTC	2400
CATCTGCCTT	ATCCAGAGTC	CGAGCTGTTC	CTTTAACGT	TTCAATAGAT	TCACCAAAT	2460
ACCAGCGACT	ACCATATACG	GCAAAATTTT	CTTTAATTC	TATAAATAAA	TTTTCGGCGT	2520
TAAATTTCTCC	TTTATTAAG	TGCAGATGAA	AATAGTCCGT	CATAATATCT	AGTACGTTTG	2580
ATGTCTCGAT	ATAATCTAAC	GAAATTTGGC	CAAAATCTCT	ATTATAGATA	AGTTGTGTCCG	2640
TTCTATCCTC	AAAACCTCCA	GTTTGAGAGT	ATTCTAACCT	TACTAGCTTG	TCTGTTAATA	2700
CAGAGATTCG	ATAAACTCT	CCCTAAAAA	TTTCAATTT	GTTTTCCTCC	TTTATGGTA	2760
GCATAAAAAC	AGAACGCACC	ATTTTGTATG	CGTTTTTCAT	TATTCTGAAT	GCAATGTTCT	2820
ATCTGTTATA	TCTATGACAA	ATAATAGTCA	ATTGAAAAA	TGCAGTGGAC	AAAATATCTT	2880
TTAACAAACC	AAGAGTTTAT	TAAAGAGTTA	TCACTTTTCA	ACTTTTCTAA	GCTTATGCAG	2940
TTGTGAAACA	AACTACTTTT	AACTATTAA	CTAAGATAGG	ATTGATAAAT	AATTTCAAAC	3000
TCTTACTAGC	AATCATACGA	TATTCAAGCT	CACGTGCTTT	TTTCCTTCCT	GCTTATTTCT	3060
TAGAACTGAA	GAACCCGGAT	CGGTATATAA	ATTATCCGGA	TCAACATAGT	CATAAGATTC	3120
ATAACAGTTG	CGTTCATTA	AGTCATCCCC	AGAGCAAGAG	CTTCATCTCG	TAATTTTCA	3180
ACATCACTAA	CCGTAGGTCG	CCATCCTTCA	ATCATATTTG	TACTTAAAGC	ATACCAAACA	3240
CTCTTAAAAA	CGGATCGGTT	TTCAAAGCT	ATTCCCATGA	TTGCATCTT	TTCTTTATCT	3300
ATATCTAAGG	ACATATGCTA	CCTCCTTTAG	ATACATTATA	CCATGTTTCT	CTGTAGCTTT	3360
TAAAAATTTT	ATTTTGTTTG	TCATATCTAA	GTTTTAGCA	CGCTTATCCT	ATTTTATAAG	3420
CCTCAAACCC	AAATATAAAA	CGCATTCTTT	TTGCTTTTTT	ACTATTGTAT	CGTATTCTAC	3480
GATAACATAC	TTTACTTTAT	TGTTTTTTTA	AATAACAGCA	GTCCCTGTT	TATCAACTAT	3540
TCGAACTACT	TTCTATTTTG	CTTCATACCC	TACATAGCGA	AAAAATATGA	AAAAGCAGAG	3600
AAGAATATCT	TAAAAAGACC	TCTTCACTGC	TAATATTAAC	ACTCATTATT	TAAACTATAT	3660
GGATTCTATC	ATCGAGTATA	CTTTTTTACT	TATTAGATAC	CTTGCTCTTC	TTTACCAAT	3720
TTTTGATCAT	ATACACGGAT	GAATGGAAGA	TAGACTAGGA	ATGCTGCAA	TGCACATACT	3780
AGAGCAACTA	ATACAGCTCG	AAGATCTGCT	GTCCTAAGA	AAGCTCCAAT	CCCTACTGGA	3840
GTTGGCCATG	GAACCTGTGC	GATAATTGGC	TTAATAAAGT	TTAGAGAATT	CGCTACGTAA	3900
TAAATAGTAG	CAGTAACCAT	TGGTGCTAAA	ATAAATGGTA	TAGCCAAGGC	TGGATTATAG	3960
ATAATAGGTA	ATCCAAAAAT	TAATGGTTCA	TTAATATTAA	ATAAGGCTGG	AACTACAGAT	4020

GCTCGTCCTA	TGCTTTAAG	CTGTTTCAGAT	TTAGAGGCAA	AAGCAATATA	TAAACATAGT	4080
CCTAAAGTTG	CACCAGAACC	ACCTGCAATT	ACAAACATAT	TAGAAAATTC	ACCTGCAACA	4140
GCGAAGTGCC	CGCCAGCAGC	ATTTTCAGCC	ATGTTAGCAA	GAGCAATTGG	ACTAACAAAT	4200
GCAAAAACAA	TGTTTCGCACC	GTGGATACCT	ACAATCCAAA	GTAGTTGAGT	CAATAGATAA	4260
ATAATCATT	AACCAATCCA	CGAATTAGTC	AGATTGGATA	CAAACCCAAA	TGGAATTGCA	4320
ATGACTTTAA	AAATATCTGT	TCCCATTGCT	ACAAGAAGAC	CGTTGATAAA	GATAACAACA	4380
AATGCAACAA	CAAATCCCGG	AACCAAAGCG	GTAATCCAC	GAGAACTCC	TTCTGGAACA	4440
GCTTCAGGCA	TTTTAATAAC	CCAATTATGT	TTAACACACA	TACGATAAAT	AAGAACAGTC	4500
ACAATTGCCA	TAATGATTGC	GGTAAAAATC	CCTGTTGTCC	CAAACGTC	GACTACATTT	4560
CCCATTGCC	ATCCATCTGC	AATTACTGCA	CCTTCTTTTA	GACTTGTAC	AGTCTTCATC	4620
ATTCCACCAT	CAAAAATGAT	TTGCGTACT	GTCATGACAA	AAGCCATCAA	GGCAAGCAAG	4680
GCACCATTAA	GAGGATTCAT	ATTGAGTTCT	TCTTCCTCTG	CATAAATTTT	TGTCAAATFCA	4740
TATGCAAGTG	ATAGAACGAA	ATAAAGAGAT	AGAGAACCCA	TAGTCGCATA	GTTTGCAACC	4800
ATGTAAGTG	ATGTGAATTT	ATCAAATGAA	GCAGAGAAAA	TATCTGCCAC	AATTGGCCAA	4860
AATGAGAAAG	CTTGTGGCAA	AATACTGAAT	ACCAAAAAACA	TTGATCCTAC	AATAGTAAAT	4920
GGTACAGCAG	CCATACCTGC	AGCCGTGATA	GCACGTACTA	CTTTAAACTG	AGCAAGTTTG	4980
CCCATTGGTC	CCATAACATG	GTTTTCAAGA	AAACCAAACA	ACCCGTTTGT	TTGATCCATA	5040
AATAGACCTC	CTTAATAAAA	CATAATAATT	TTTACTTTCT	AAAGACTAGT	TTCAAATACA	5100
AATTATACTA	GATCAGGATT	ATAAACTAAG	TGAGTTCTTT	TCCAATGGA	CAAATTGTTG	5160
ATAAGCCTTA	TCTGTTGTT	TATAAATTTT	TTAATTTCTT	CTAATGTCTA	ACAAACTCAG	5220
AACTAAACCT	AATAGAAGAA	CTACAAAAAC	AAATAAACGT	GCTACTTGGT	TATTTTCAAA	5280
AATCGAAAA	AGATTCCTTA	ACCAACTTGT	CCAAGTTAAA	ACAAGTAATC	CTATTGAAAT	5340
AAGCATTTGT	ATTCTAACAA	ACATTAGTGT	TATTCCCAAC	TTTTCTTCC	TATTTCCAFA	5400
AAGTTTAAAT	TGTTCAACAG	TTGCTAAAAT	AGAAAATACT	ATGAGCATAA	TGGGGAAAAAT	5460
AATAATAGGC	GAGGGACTAA	TAAACTGACT	CAAAGCCAA	TAAATATTC	CAAAAAAGAA	5520
GAGTGCTATT	GAATAACGTA	GAAGAAGATA	TCGATTGAAA	AAAGTATTAG	TTAGAGCCAT	5580
CTCTCGACGT	TGTTGTTCAA	TCTTTTGTG	TTCTTTTTTA	TCCATATCAT	TFCTCCTTA	5640
TATAACAACA	CATATTTAGT	TAACTTTCTT	ATAAAGAGCT	AACATTTCTT	TTGCTACTTC	5700
TAATAATGTC	ATAGTGGTCA	TTAAATGATC	TTGAGCATGT	ACCATGATAA	TTTCAATTTT	5760
AATTTCCACT	CCACTTGCCT	ATTCTTGCAA	GAGTTTGGTT	TGTGCATGAT	GCGCTTCAAG	5820

AATTATCTCA TTTGATTGAT TTAATTTACT TTCTGCATCA TCAAAACTAC CTTCTCTCAT 5880
 TTTTGCAAAAT GCTTCATGTA TTTCTGACCT TGCATTTCCC GAATGCAGGA TAATTTCAAA 5940
 TGCTGCAACC TGCAGTTCCT CTGATTTCAT ATAAACCTCC TATTTTATCT TCTCAAATAT 6000
 GTTAATAAAA TCTTCAAAGT TATTGCAAGA TATTAGCTGA TTTTGCAATT CATCATTCTC 6060
 TGTCAGAGAG ACTATCTTTT TAGTCACAGT TGCCAAACCT TCGTTCCCAT ATATTGATGG 6120
 AGATAGAAGA AATACTAGCT GGACATGTGA ACTTTGATTA TCCCAGAGTA ACGAATCTTT 6180
 ACAAATTGCA ACCGAAACCT TTCCCTCTGT ACCAAAGGGC TGAATAGGAT GCGGAACCTGC 6240
 AATTTTTTCA GAAAAACAA CTGAACCTAA TTCTTCGCGC TGTTTAATTC CATAAAGTAA 6300
 AGATTGTTCA AACTCATTG ATTCACCAAC AGATAAACTC TCAACCATCT TTCAAGTAA 6360
 ATTTACCTTG TCTGATTTCAG TACATATTAA AAGTTTCTCT TTAATAAAAT ACTGTCTAAA 6420
 GCCGTGTTTT TCAAATTTGT TAATCTTTGA TGATTGTACA TAACTAGAAA CTTGCATCTA 6480
 ATCCATAGCT TTTCTAATCA TTTCCATCTC ATCACTCTTA AGAAACACAC TAACTTTAAA 6540
 AACTGGGATT TGAAAATATA GATTTGATAA ATCAATAGCT GACACTATAA AATCTATTCC 6600
 TTTAAGTTTT TCTTGATTCA ATTCATAGTA GCCTATTACA TCAACAACCTT CTAICTCGCTT 6660
 CCCAACTCC GTTTCCAAAC GATTTCTTAA CATTTGGGCT GCACCAAATC CTGTTGCACA 6720
 AATAGCAAGA ATATTAAACT TAGTACTCTC TTTGCTACGT TCCATAGCAG CTAAAAAGTG 6780
 AAGACTTACA TATGCTACTT CATCATCTGA TATTGTCCAC TCCAAGAACT TGTCATATT 6840
 TGCAAGAATT TCTCTAGTCA TAAAGAATAT ATCACTATAA TTCTGTTTAA TTTTCATCTAC 6900
 CAAAGGGTTA TTTAAGGTAA TCCGGCTTTC TAAACGFACT TGTAGTGTC TTAGATGAGT 6960
 TATCAATCCT TCAATTAGTT GGAAATCTGA AGAAAAGTTA TACATATCAT CTAATCCTAA 7020
 ATTCTGAAAT GTTTTAAATA AAGATTTTTT TAAAACCTCT TCAGAAATAT TCTTCTGATT 7080
 TTTTGTGACAT TGTTGACTCT TAGCTAACAA ATGCAAAGTA ATGTAGTCTA TTTCTGAAAC 7140
 TGGAAATTC TGATTTGTTA CTTCTCTTAC TTTAGAAAGA ATTCTTTGGG CAACCTTTCT 7200
 CTCTATTGCA TCATCAGTCA TCTGACAGTC TATATTTTTT ATTTCAAATC CGGATTTTAA 7260
 ACGAATCACA GACAATGCTA TGTGAACTAC TAAATTCCTGT AGTACAAAAT CAGATAGTTT 7320
 TAGGTTGGCC TCTTGGCATT CATCCAAAAC AATTCTAGCA AATTCTTCTA ATGGAACAGT 7380
 TTGATCAAAA AAGTTAAATT TTACATAGCA ATGTATTGTT TTAATAAATT GATTCTCTAG 7440
 GAAATAATTT ATGATAAAC GTCGTTTATC ACGTTCCTCG CCTGAGACAT AAACCTCTTT 7500
 ATTCCGCCCTA CTCTCAATGG ACAAATATA CTCTGATAAC ATCACTCGTA TCTTTCTGAA 7560

ATCATGAGAT	AATGTTGAAC	GACTAACGTA	AAGTTCATCA	GCTAAATCAT	CAAAAAGAAC	7620
TGGAACCTGC	TCAAATAATA	ATTTATTTAA	GATAAATACT	AAACGATCAT	CACCTTTTGA	7680
AACCGCAGTT	TTCGTATAGT	CTTCTTCCAG	TTCATAAGTT	TGTCTAAACT	CCTGGTAAGC	7740
GCCTTGATTC	TCAAAAAATA	TTTGATACCC	TTGACCTTGT	TTTGAAATCA	ACCGGACTCC	7800
TTGAATAATC	ATTGTCCTCT	CAATTAATTT	CAGTACATTA	CGGACAGTTC	TATCTGAACA	7860
GGATAAATAT	TCTGCCAGTT	CTTTGCTTGT	AACAAAACGT	TCCTTATTTT	TTATTAATAA	7920
TTGAAGGATA	TCTTTCTCTT	TAATGTTTAA	CACATTCATT	CCCTCCTAAA	ACGTATGTTT	7980
TCATATATTG	AAGCATATTA	TACACTTAAA	TCAGTTTATA	TCAAACCTCA	AACAATTTAT	8040
CTTAACCTAA	ATATTTATTG	ACATTTTCATG	TGTTTCATCA	ATATTCCTCA	GAATCAAATT	8100
AGCCATTTTT	TCAATTCCCA	TTGGAATAGG	AATATAGGCT	TGAGGAGGTA	TTTGTACAAC	8160
TGGTTTTCCCT	GCTTTAGAAC	CAGCCTCTTC	AAATTGCTTA	AAGTACATTT	TTGTTTGAGG	8220
ACTGACAAGA	TACAAATCAA	AAGCTGCTGC	TGCGATAGCT	TTCCCTCCTT	CAGTAGCACT	8280
AATAGCATCA	ACTACAATAT	CTTCCCTTTT	TCCTTTTAGA	AACTCTGTTG	TTTTCTGTGC	8340
CATAAGTGAT	GAAGACATTC	CTGCTGCACA	AATAATTTAA	GCTTTTGCCA	TAATATTTTC	8400
TCCTTTTCTT	AAATCCAATC	AAAGCTGTGC	TAAGTTGGCT	TATTTGTAT	CTATTTTAT	8460
TATAAAATAA	AGCGTTTCCA	ATGACAATTC	CCTCATTTTC	CTAAATGATA	TGGAAAAAAA	8520
TTATTTATAC	TTCAATTTAT	AAAATAAAAT	TATTCCTGAG	AGTAGAAATG	AAACACTATT	8580
TGCTAAAATC	AAAGGCAAGT	CTCCTATACG	AATACCATGA	GCAAGCCACA	ATGCAATACC	8640
AATAACTTGC	ATAACATACA	TACCTAGAGC	AATAGATCCT	GTGTCCTTTG	TCTTAACCTAC	8700
ACGAAAAACT	TGTGGTAAAA	ATGCAAATGT	TGTTAAAATT	GCTGCAATAC	TTCCAATCAT	8760
ATGTCACCTC	AATATGCTAA	ACAAACTGAG	AATAATCTCA	GTTTGTTTAT	ACTATTCTAC	8820
TGATTCACCG	TTAGATGAAA	TAACCTCCTT	ATACCAGCCA	AAAGATTTTT	TCGGGGAACG	8880
ATTATAACTT	CCCTTCCCAT	TATCATCTTT	ATCTACATAA	ATAAAGCCAT	AACGTTTCCG	8940
CATTTACCG	GTACCAGCTG	AAACCAAATC	AATACATCCC	CATGGAGTAT	AACCCATTAA	9000
ATCAACACCA	TCTTCAACTA	CAGCCTTTTT	CATTTACCGA	ATATGGGCAC	CTAGATATTC	9060
AATTCATAA	TCATCATGTA	CCATACCATC	TGCTGCAACT	TGATCTATAG	CTCCAAAACC	9120
ATTTTCAACA	ATAAAGAGTG	GTAAGTGATA	GTGGTCTGTA	AACCAATTTA	ACGCATAACG	9180
CAAACCTTCT	GGATCAATTT	GCCACTCCCA	TTCAGAAGCC	TTAACATAAT	TATTTTTCAC	9240
TAAATCTTCT	GTTTCAAGAT	AATCAAAATA	AGGATTATTT	TCACGATGAG	AGTCGATAGC	9300
AAAGGACATA	TAGTAACTGA	AACCAATGTA	ATCTACAGTC	CCACCAAGTA	AATCTTCTTT	9360

ATCCTGGGCA	GTAAAATCAA	CTGAAATACC	TTTTTCGTTCC	CAATACTTGA	AAATATGCTC	9420
AGGATATTTA	CCTAAAACAT	GCACATCAGC	AAAATAATAA	CGCTTCTGCA	TAGCTTTCAT	9480
TGCCATTAAG	ATATCCTTAG	GATTGCAAGT	AACTGGATAA	ATTGGACACA	TCGCAATCAT	9540
ACAACCTATT	TGAAAATCTG	GATTAATCTC	ATGACCAATT	TTTACAGCTC	GTGCAGAAGC	9600
AACTAATTCG	TAATGTGCTG	CTTGATACAT	AATTGCTTCT	CTATTATCAC	CTTCCTCATA	9660
TACAATACCT	GAGTTAGTAA	ATGGTGCAAA	ATCTTCCTGA	TAATTCGCTT	GATTATTGAT	9720
TTCATTGAAA	GTCATCCAAT	ATTTAACCTT	ATCTTTGTAA	CGTTTAAATA	CGACTTCTGC	9780
AAAACGAGCA	AAGAAATCAA	TCAATTCCTT	ATTTTCCCAA	CCACCATATT	CGGTCACTAA	9840
GTGATAAGGC	ATTTCAAAT	GAGATAGAGT	GATGACAGGT	TCAATACCAT	TCTTTAAGCA	9900
TTCATCAAAA	AGATTATCAT	AAAACGTAA	TCCTTCTTCA	TTCGGCTCTA	ACTCATCACC	9960
TTTTGGAAAG	ATACGTGTCC	ATGCAATAGA	GGTACGGAAG	CACTTGAATC	CCATTTCAGC	10020
AAAAAGTGCT	ATATCTTCTT	TATAACGGTG	ATAAAAATCT	ATCGCCTCAT	GATTTGGATA	10080
ATATTTACCC	TCTAAAACCT	CCAAAGTAAT	TTCACGAGCT	ACTCCATGAC	GACCAGCAGT	10140
CATAACATCA	GCAACACTAA	TTCCCTTGCC	ACCTTCTTGC	CATCCACCTT	CAAGTTGATG	10200
AGCAGCAACA	GCACCACCCC	ATAAAAATCC	ATCTTTAAAA	GTAGTCATCT	TTTTTCCTCC	10260
TGACTTTGAT	ACTCTTATTA	TAAACCTTAA	ACCAAAAGAT	GAAAACGCAT	TCTTTTTCCT	10320
TATTGTTAAG	GAAAGAAGTA	ATTTTAAATG	GAAATAGAAC	AATATCTTCT	TGTATTTCTCG	10380
TAATGATATC	TTTACGATTT	TCAATACTTT	CAAACACTAA	AAACTCTCAC	AATAAATCTA	10440
ATTCCCTGTG	TCTATAAACG	ACTTATCGCT	TTCTGGCATC	CCAGAATCAT	CTTCTATATA	10500
ACGTTCAACT	TGCATCTGCA	AGTGATATTT	TTTTCTTAAA	TCTAAGATTT	TCTGCATTGT	10560
CTTTGATTGA	TAATGTTTAT	CTAAAGTTTC	TTGATTTATC	CACTGATCAA	TAAGGAGAAT	10620
AGTTCCCTCT	TTTTCAATTG	GTAAAAAATA	TTCGTATTTT	AAGTTACCTT	TTTGATTTCT	10680
AATTTCTTTA	ACAAGGCCAC	TATCAAGCAT	TTCTCTTGCA	AACTTTATTG	CACTATCTCC	10740
ATCACCTTTA	TAATATACAT	GAATAGTCAA	TGTCATCTTA	TATCCTCCAA	AATCATCCTT	10800
CAATTTTAAA	AAAACAAGTT	TAGATGAGGA	TCTAAACTTG	TTTTTTATGA	ACTAATTATC	10860
TAACGTTTCG	CCATTACTTT	CAATCACTTC	TTTATACCAA	TAAAATGATT	TTTTCTTATA	10920
GCGATTTATA	GTCAATTGAA	ACAAGAGCAG	GACAAAAGAG	CCTCATAAAA	GGTATTGCAA	10980
CTTGGAATA	CCTTTTGTAG	GTGCTTTTGT	ATATGAGCCC	ATGTTTTTCTC	AATAGGATTG	11040
TACTCAGGTG	AGTAGGGAGG	AAGAGGTAAA	AGTTTATACC	CAAACCTTTC	ACACAAGAGT	11100

838

TCTAGCTTCC CCATTCTATG GAATCTTGCA TTATCCATAA TAATAACCGA TGGTGTGGTT 11160
 AATGTTGGTA AGAGAACTT CTGAAACCAA GCTTCAAAAA AGTCGCTCGT CATCGTCTCT 11220
 TCGTAAGTCA TTGGAGCGAT TAACTCACCA TTTGTTAGAC CTGCAACCAA AGAAATCCTC 11280
 TGATATCTTC TTCCAGATAC TTT 11303

(2) INFORMATION FOR SEQ ID NO: 116:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 3112 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 116:

CCTTAGATTT CCACTTGCCA GAGGAATTGA TTGCCCAAAC GCCCCTTGAA AAACGTGATG 60
 CCTCCAAACT CCTCATCGTC AACCGTGAGA CAGGAGAAAT GCAAGATAAA CATTTCCTACT 120
 CTATTATTGA TATGCTGGAA CCTGGTGATG CCCTTGTTCAT GAACGACACC CGAGTTCTCC 180
 CTGCCCGCCT CTATGGTCAA AAAGTGGAGA CAGGAGGTCA TGTGGAACTT CTCCTCCTTA 240
 AGAACACTAG TGGAGACGAG TGGGAAGTTC TGGCTAAACC TGCCAAACGC CTCAAGGTCG 300
 GTACTCGTAT CAGCTTTGGT GATGGCCGCC TCAGCGCTGT CGTTACAGAA GAATTGACCC 360
 ACGGGGGACG CATTGTCCGC TTTGAATACC AAGGAATTTT CCTAGAAGTC TTGGAAAGTC 420
 TGGGAGAAAT GCCTCTGCCA CCTTATATCC ACGAAAAATT AGATGACCGT GAACGTTATC 480
 AAACCGTCTA CGCCAAGGAA AGTGGCTCTG CTGCAGCACC GACTGCTGGT CTTCACTTCA 540
 CCAAAGAACT GCTGGCAGAA ATCCAAGCTA AGGGTGTTC A TCTAGTCTAT CTGACTCTCC 600
 ATGTCGGACT CGGAACCTTT AGACCTGTTT CTGTGGATAA TCTGGACGAA CACGAAATGC 660
 ACTCAGAGTT CTATCAACTT TCTGAGGAAG CTGCTGCCAC CCTTCGCTCT GTCAAAAAAA 720
 ATGGTGGTCG TGTCATCGCT GTCGGAACCA CTTCTATCCG CACCTTGGAA ACTATTGGTT 780
 CCAAGTTTGA TGGGCAAATC CAAGCAGATT CTGGTTGGAC CAATATCTTT ATCAAACCTG 840
 GGTATGAGTG GAAGGTCGTG GATGCCTTCT CAACCAACTT CCACCTGCCA AAATCAACTC 900
 TGGTCATGTT GGTTCCTGCC TTTGCAGGCC GTGAATTAGT CTTAGATGCC TACCACCATT 960
 CCATCCAAGA ACACTACCGC TTCTTCAGTT TTGGTGACGC CATGTTTATT TATTGAGAAA 1020
 GAATTTCTCT AAATCTTCTA ATACCAATAA ATCGCTAAGA TATTATTTCA AAGAACATCT 1080
 ACAATTGAAA CTCTAGCTAG CTGTAGAAGA GGCCTAGTAC ATTGAAATTA AAATGCTTCC 1140
 CCCTAGCTTC GAAAATATTG CCATAGATTG CGTTGACTCT CCAAATGAT TCATCTATAT 1200

TTTATTTTCAG CTTCCTATAC TTTCTTCGCT GTTTGTAAT CAAAATGCAA GACACATGAG 1260
 TAGCACCATATA TTTGTTACTC TTATCTGTCC TCTCAAGAGA CTATTATGAG TTATTTTCAGA 1320
 ATCATTTCACT ACTTTGACCC TGACTCTCCT TAGTCTCAAA ATCAAAGACT TATACTCTTC 1380
 AAAAATCTCT TCAAACCGCG TCAACGTAC CTTGGATTAT ATATGTGatC TGaCTTCGTC 1440
 AGTTCATCT ACAACCTCAA AGCAGTACTT TGAGCAACCT GCGACTAGTT TTCTAGTTTG 1500
 CTCTTTGATT TTCATTGAGT ATTAACAACA AAGTGAACAA ATCTGAATTC TAATGTACAG 1560
 AAGACTAGGC TTGTTCACTT TTTTATAGTC GCTATAAGAT GACCTTATCT ATAGCTTTTT 1620
 ATATATAAAT ATATATTCAG ACATACTATT ATCAATTTTG TCGCAGGGAG GAATCTGTTA 1680
 ACGCACCCAT TCACCATTAT CATTGACTCT ATAGCCATCT ATACTTGTAT TGACCCTAA 1740
 CTCACCCGAT GTATTTACAT AATACCATTT ACCACCAACT TGGAACCATT GATTGACTTT 1800
 CATAGAACCG TTGCTGTTGA GGTAGTACCA TGAACATTA ACTTGTACCC AACCTGTTGC 1860
 CATGGAACCA TCAGTATTAT AAAAATACCA CATACCATTT TCTTGTTC AGTCTGTGT 1920
 TGGAGCAACT GCTTTAGCTG GTTCTACTGC TACATCTGTT CCTTGGTTAG ATGTAACAGA 1980
 TACAGGATAC GAAGGAATAG ATGATTGCTC AGGAACAACA ACTTTTTCAG GTTCTCTCGT 2040
 CCCTCTCCTT ATACGTCTTT TTACCATCTC TTTAGTAATT TGACGAGAAG TAGTTTCTTC 2100
 AATGTTTCCA TCACGTTTCA CTACAGTATA GATGTAGTA AGAGTAATTT ACCAATTTCT 2160
 CCTACTTCTT CTACTTCTTG ACTTTTATCA AGAGTTGGGC CATCGAGATA TTCTGTTTCG 2220
 ATTGGAATTT CTGGACAAG AACTTGGGGC TTGGTTCTTT TTTTAACAAC TCTTGTTTGA 2280
 GAGTCTTTTT TTTGACTTAA AGTACTCTCA GTTACTTGTC CACTCTTCC ATCTACATTA 2340
 TAAGTTATCG TTGTAACGT TTTCCCATTC TTTCTAGAG TAATCTCTTG CTCCTGTCC 2400
 GCAGAAAGGT CATGTCTGC TTCATATTTA GTAGCAAATG GAACAAGAAC TTCTTCAACC 2460
 TTGCTTTTAG CTGGAACCTT GATAACTGTA TCCGTGGCTT CTTTCTATC AACAGTAACC 2520
 TGTTCGGTAA CATAACCAGT CTCTGGATTA ACATCGTAGG TCCTTGTCGT AGTTACATAG 2580
 CCATCCTCTC CATCAATTGT AACAGGATTT TCACTACGGT CTTTGTGTTT ATCTTTTTCA 2640
 TAACGAATTC GCGTACTTGA AATTTTCTTG GTTACTACCT TAGGTTTAGT CGTACTTTT 2700
 ACAATAATAT CCCCATTTGTC AGCGTCATCA TACTCTATTC CCTCTTCTTT ATCTCTAGTA 2760
 TCATCTCTGA CATATTGAAT CCCATCAGCA GCATGAACAA AACTTGATTT CAGATTCCTC 2820
 CTAAAAATAA AGTTAGCCCG ATTACCGCAG AACCAAAAAT CTTTCCGAGT TTACGTATTG 2880
 CATAGCGCTT ATTAGTATTA GATTTTGCCA TTACATCCTA CTTCTAGTAT AGCATCTTTT 2940

840

CTATCAAACG TTAAACAATA TACGTTATAT ATAAAATAGA CTTAGAATGA TATATTGATT 3000
 ATTGAACTAA CACTTTAACT ATATCGTAAT CAATCTCATA TATAAAGGAT TGCAGACATC 3060
 TTATCTAAAT ACATGCGAAT ATATTTAGAT ACAAACATTC CAACTTGATA AT 3112

(2) INFORMATION FOR SEQ ID NO: 117:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 4327 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 117:

CCCAAAAATC TCTTCAAACC ACGTCAGCTT CGCCTTGCCG TAGTATGGTT ACTGACTTCC 60
 TCAGTTCTAT CCACAACCTC AAAACAGTGT TTTGAGCATC ATGCgGCTAG CTTCTTAGTT 120
 TGCTCTTTGA TTTTCATGTA GTATAAAAAC AGATGAGTTT CTGTTTTCTT TTTATGGACT 180
 ATAAATGTTC AGCTGAAACT ACTTTCAAGG ACATTATTAT ATAAAAGAAT TTTTGAAC 240
 TAAAATCTAC TATATTACAC TATATTGAAA GCGTTTTAAA AATGAGGTAT AATAAATTTA 300
 CTAACGCTTA TAAAAAGTGA TAGAATCTAT TTTTATGTAT ATTTAAAGAT AGATTGCTGT 360
 AAAAATAGTA GTAGCTATGC GAAATAACAG ATAGAGAGAA GGGATTGAAG CTTAGAAAAG 420
 GGAATAATA TGATATTTAA GGCATTCAAG ACAAAAAGC AGAGAAAAG ACAAGTTGAA 480
 CTACTTTTGA CAGTTTTTTT CGACAGTTT CTGATTGATT TATTTCTTCA CTTATTTGGG 540
 ATGTGCCCT TTAAGCTGGA TAAGATTCTG ATGTGAGCT TGATTATAT TCCATTAT 600
 TCTACAAGTA TTTATGCTTA TGAAAAGCTA TTTGAAAAG TGTTCGATAA GGATTGAGCA 660
 GGAAGTATGG TGTAAATAGC ATAGGCTGAT GTCCATCATT TGCTTATAAA GAGATATTT 720
 AGTTAATTG CAGCGGTGTC CTGGTAGATA AACTAGATTG GCAGGAGTCT GATTGGAGAA 780
 AGGAGAGGGG AAAATGGCA CCAATTTGAG ATAGTTTGTT TAGTTCATTT TTGTCATTIA 840
 AATGAACTGT AGTAAAAGAA AGTTAATAAA AGACAAACTA AGTGCATTTT CTGGAGTAAA 900
 TGTCTTATTT CAGAAATCGG GATATAGATA TAGAGAGGAT CAGTATGAAT CGGAGTGTTC 960
 AAGAACGTAA GTGTCGTAT AGCATTAGGA AACTATCGGT AGGAGCGGTT TCTATGATTG 1020
 TAGGAGCAGT GGTATTTGGA ACGTCTCCTG TTTTAGCTCA AGAAGGGGCA AGTGAGCAAC 1080
 CTCTGGCAAA TGAAACTCAA CTTTCGGGGG AGAGCTCAAC CCTAACTGAT ACAGAAAAGA 1140
 GCCAGCCTTC TTCAGAGACT GAACTTTCTG GCAATAAGCA AGAACAAGAA AGGAAAGATA 1200
 AGCAAGAAGA AAAAATTCCTA AGAGATTACT ATGCACGAGA TTTGGAAAAT GTCGAAACAG 1260

841

TGATAGAAAA	AGAAGATGTT	GAAACCAATG	CTTCAAATGG	TCAGAGAGTT	GATTTATCAA	1320
GTGAACTAGA	TAAACTAAAG	AAACTTGAAA	ACGCAACAGT	TCACATGGAG	TPTAAGCCAG	1380
ATGCCAAGGC	CCCAGCATTC	TATAATCTCT	TTTCTGTGTC	AAGTGCTACT	AAAAAAGATG	1440
AGTACTTCAC	TATGGCAGTT	TACAATAATA	CTGCTACTCT	AGAGGGGCGT	GGTTCGGATG	1500
GGAAACAGTT	TTACAATAAT	TACAACGATG	CACCCTTAAA	AGTTAAACCA	GGTCAGTGGA	1560
ATTCTGTGAC	TTTCACAGTT	GAAAAACCGA	CAGCAGAACT	ACCTAAAGGC	CGAGTGCGCC	1620
TCTACGTAAA	CGGGGTATTA	TCTCGAACAA	GTCTGAGATC	TGGCAATTTC	ATTAAAGATA	1680
TGCCAGATGT	AACGCATGTG	CAAATCGGAG	CAACCAAGCG	TGCCAACAAAT	ACGGTTTGGG	1740
GGTCAAATCT	ACAGATTCGG	AATCTCACTG	TGTATAATCG	TGCTTTAACA	CCAGAAGAGG	1800
TACAAAAACG	TAGTCAACTT	TTTAAACGCT	CAGATTTAGA	AAAAAACTA	CCTGAAGGAG	1860
CGGCTTTAAC	AGAGAAAACG	GACATATTCG	AAAGCGGGCG	TAACGGTAAC	CCAAATAAAG	1920
ATGGAATCAA	GAGTTATCGT	ATTCCAGCAC	TTCTCAAGAC	AGATAAAGGA	ACTTTGATCG	1980
CAGGTGCAGA	TGAACGCCGT	CTCCATTCGA	GTGACTGGGG	TGATATCGGT	ATGGTCATCA	2040
GACGTAGTGA	AGATAATGGT	AAAACCTGGG	GTGACCGAGT	AACCATTACC	AACTTACGTG	2100
ACAATCCAAA	AGCTTCTGAC	CCATCGATCG	GTTCAACAGT	GAATATCGAT	ATGGTGTGG	2160
TTCAAGATCC	TGAAACCAA	CGAATCTTTT	CTATCTATGA	CATGTTCCCA	GAAGGAAGG	2220
GAATCTTTGG	AATGTCTTCA	CAAAAAGAAG	AAGCCTACAA	AAAAATCGAT	GGAAAAACCT	2280
ATCAAATCCT	CTACCGTGAA	GGAGAAAAGG	GAGCTTATAC	CATTCGAGAA	AATGGTACTG	2340
TCTATACACC	AGATGGTAAG	GCGACAGACT	ATCGCGTTGT	TGTAGATCCT	GTAAACCAG	2400
CCTATAGCGA	CAAGGGTGAT	CTATACAAGG	GTGACCAATT	ACTAGGAAAT	ATCTACTTCA	2460
CAACAAACAA	AACTTCTCCA	TTTAGAATTG	CCAAGGATAG	CTATCTATGG	ATGTCCTACA	2520
GTGATGACGA	CGGGAAGACA	TGGTCAGCTC	CTCAAGATAT	TACTCCGATG	GTCAAAGCCG	2580
ATTGGATGAA	ATTCTTGGGT	GTAGGTCCTG	GAACAGGAAT	TGTACTIONCG	AATGGGCCTC	2640
ACAAGGGACG	GATTTTGATA	CCGGTTTATA	CGACTAATAA	TGTATCTCAC	TTAGATGGCT	2700
CGCAATCTTC	TCGTGTCATC	TATTCAGATG	ATCATGGAAA	AACTTGGCAT	GCTGGAGAAG	2760
CGGTCAACGA	TAACCGTCAG	GTAGACGGTC	AAAAGATCCA	CTCTTCTACG	ATGAACAATA	2820
GACGTGCGCA	AAATACAGAA	TCAACGGTGG	TACAACATAA	CAATGGAGAT	GTAAACTCT	2880
TTATGCGTGG	TTTGACTGGA	GATCTTCAGG	TTGCTACAAG	TAAAGACGGA	GGAGTGACTT	2940
GGGAGAAGGA	TATCAAACGT	TATCCACAGG	TTAAAGATGT	CTATGTTCAA	ATGTCTGCTA	3000

842

TCCATACGAT GCACGAAGGA AAAGAATACA TCATCCTCAG TAATGCAGGT GGACCGAAAC 3060
 GTGAAAATGG GATGGTCCAC TTGGCACGTG TCGAAGAAAA TGGTGAGTTG ACTTGGCTCA 3120
 AACACAATCC AATTCAAAAA GGAGAGTTTG CCTATAATTC GCTCCAAGAA TTAGGAAATG 3180
 GGGAGTATGG CATCTTGTAT GAACATACTG AAAAAGGACA AAATGCCTAT ACCCTATCAT 3240
 TTAGAAAATT TAATTGGGAA TTTTGTAGCA AAAATCTGAT TTCTCCTACC GAAGCGAACT 3300
 AGAGAGATGG GCAAAGGAGA GATGGGCAAA GGAGTTATTG GCTTGGAGTT CGACTCAGAA 3360
 GTATTGGTCA ACAAGGCTCC AACCCCTCAA TTGGCAAATG GTAAAACAGC GACTTTCCTA 3420
 ACCCAGTATG ATAGCAAGAC CTTGTTGTTT GCAGTAGATA AGGAAGATAT CGGACAGGAA 3480
 ATTATGGTA TAGCTAAAGG AAGCATCGAA AGTATGCATA ATCTTCCTGT AAATCTAGCA 3540
 GGTGCCAGAG TTCCTGGCGG AGTAAATGGT AGCAAAGCAG CGGTGCATGA AGTTCAGAA 3600
 TTTACAGGGG GAGTTAATGG TACAGAGCCA GCTGTTCATG AAATCGCAGA GTATAAGGGA 3660
 TCTGATTCGC TTGTAACCTCT TACTACAAA AAAGATTATA CTTACAAAGC TCCTCTTGCT 3720
 CAGCAGGCAC TTCCTGAAAC AGGAAACAAG GAGAGTGACC TCCTAGCTTC ACTAGGACTA 3780
 ACAGCTTCT TCCTTGGTCT GTTTACGCTA GGGAAAAAGA GAGAACAATA AGAGAAGAAT 3840
 TCTAAACATT TGATTTTGTG AAAATGGCTC TTTGTCAACT GTAGTGGGTT GAAGTCAGCT 3900
 AAGCTCGAGA AAGGACAAAT TTTGTCCCTT CTTTTTTGAT ATTCAGAGCG ATAAAAATCC 3960
 GTTTTTTGAA GTTTTCAAAG TTCCGAAAAC CAAAGGCATT GCGCTTGATA AGTTTGATGA 4020
 GATTATGGT CGCTTCCAAT TTGGCGTTAG AATAGTGTAG TTGAAGGGCG TTGACGATTT 4080
 TCTCTTTGTC CTTTAGAAAG GTTTTAAAGA CAGTCTGAAA AAGAGGATGA ACCTGCTTTA 4140
 GATTGTCCCT AATGAGTCCG AAAAATTTCT CCGGTTCCCT ATTCTGAAAG TGAAACAGCA 4200
 AGAGTTGATA GAGCTGATAG TGATGTTTCA AGTCTTGTGA ATAGCTCAA AGCTTGTTTA 4260
 AAATCTCTTT ATTGGTTAAA TGCATACGAA AAGTAGGGCG ATAAAAATGT TTATCGCTGA 4320
 GTTTACG 4327

(2) INFORMATION FOR SEQ ID NO: 118:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 3521 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 118:

CTCTGGCCCT GCCACTCCAA CGTTTGTCA GGGTGCTTTT TTCATAAAGG AGTTCTTATG 60

843

TTAGATATCA AACGTATTCG TACAGATTTT GAAGCTGTCG CAGAAAAATT AGCTACACGT	120
GGTGTAGATG CTGCTGTCTT GAATGAAATG AAAGAAATCG ATGCTAAACG TCGTAACATC	180
TTGGTCAAGG TTGAAACTCT CAAAGCAGAA CGTAACACAG TTTCTGCTGA GATTGCCCAA	240
GCTAAGCGCA ACAAGGAAAA TACAGATGAC AAGATTGCTG CCATGCAAAA TCTATCTGCT	300
GAGGTTAAAG CCTTGGATGC TGAATTGGCA GAAATCGATG CTAAATGAC AGAATTTACA	360
ACGACTCTTC CAAATATCCC AGCTGACAGC GTTCCTGTTG GGGCTGACGA AGACGACAAT	420
GTGGAAGTTC GCCGTGGGG TACTCCACGC GAGTTTGA CTGAACTAA AGCTCACTGG	480
GATCTCGGTG AAGACCTTGG TATCCTGAC TGGGAACGCG GTGGTAAGGT AACAGGCCT	540
CGCTTCCTCT TCTATAAAGG CCTCGGTGCT CGTTTGGAAC GTGCTATCTA CAACTTTATG	600
TTGGATGAAC ATGGAAAAGA AGGCTATACT GAAGTCATCA CACCTTACAT AGTCAACCAT	660
GATTCTATGT TTGGTACTGG TCAGTATCCA AAATTTAAGG AAGATACTTT TGAATCAGC	720
GATACCAACT TTGTCTTGAT TCCAACGCT GAAGTTCCTC TGACAAACTA CTACCCTGAT	780
GAAATCTTAG ACGGCAAAGA TCTTCCAATC TACTTCACTG CCATGAGTCC GTCATTCCTG	840
TCTGAGGCTG GTTCTGCCGG TCGTGATACG CGTGGCTTGA TCCGTTTGA CCAATCCAC	900
AAGGTGAAA TGGTCAAATT TGCCAAACCA GAAGAATCTT ACGAAGAATT GGAAAAATG	960
ACAGCCAACG CTGAAAACAT TCTTCAAAAA CTCAACCTTC CATACCGTGT CGTTGCTCTC	1020
TCTACTGGAG ATATGGGCTT CTCAGCTGCG AAGACTTACG ACTTGGAAGT GTGGATTCCA	1080
GCACAAAACA ATTACCGTGA AATCTCAAGC TGTTCAAACA CAGAAGATTT CCAAGCCGT	1140
CGTGCCCAA TCCGTTACCG TGATGAAGCA GATGCAAGG TGAAACTCCT TCATACCTTG	1200
AACGGTTCTG GACTTGACGT TGGACGTACA GTGGCTGCAA TTCTTGAAAA TTACCAAAT	1260
GAAGATGGTT CTGTGACCAT CCCAGAAGCA CTTGTFCCAT ACATGGGTGG AGCTGAAGTC	1320
ATCAAACCAT AAAAAATAAG GTTTAGCTAT TTCTAGCTAG ACCTTTTTTC GTAACCAAAT	1380
CAGATAAGCA CCTAGTACAA AGAATAAAAT AGTTAGGCAT ATAATGGTTT CAGCCAATAC	1440
CAGGTAATCC AGAAATGGAA GTTTCAAAAT TCCCTGAGCC ATCTTGAGCG AGGTCGCTGT	1500
GATAATGGTT GGAAGGTGA GGGCTGAGAA GGCTGGTTGA AAACCTTGT TTAATATGTT	1560
GGGCAGACGA GTTAAACAA AGAAAAAGAA GGATTGAGAA GCCAAAATCA TGACAATCAA	1620
GACCCAAGTC GGCAGGCTGG TTCCTCCTAC TCGAACTAGA GAAGCCAAGA GTAGAGAGAA	1680
AGGAGCACAG TAGATTCCTT CTTGTCCAAG CAAGGCTAGT GGGAGTGGAT GTTCTTTAA	1740
ATCGCTATAA ATAAGGGGAT AGAGATAGAA GGTCAAGAGA AAACCAAAC TCAAGTCCG	1800

844

ATAGGCAATT TCGATAATAC CTACCAGAGG ATAGGTCAAG GCAGCCACTG CTATCCCCAC 1860
 ATAGAGAACC GTCCAGCTTG GAGTGGCATG AACCTCCGC CCTGGACAAG CAAACTTGAT 1920
 GGTAAAACCA GCAATCAAGG TCAAATCCAA GAGAAATGAA AACCAACAAA TCCCTTGTGC 1980
 TACCAAAGGA AGATAAGAGA ATACGCGAAA GACATAGGTC GATAAAATCA TCCCAGCCAT 2040
 AGGAAAGGTT GCCATTCTG ACAAAGAGG GGGCTTGGTC AATTCTTGCT TGGTTCTTT 2100
 CCAATTAAG AGATGCAGAA TTAGAAAGTA AATCCATAAA ACCAAACCAA TCAGACTAAA 2160
 AAGATGGGAT AGAACCGGCA ACGTATCTAA AATAAGATTT CCAGCTCCTG CCAAACCTAG 2220
 CAAACAACCT GAAAATACTA AGGGGAGTTT TTTCATCCTA ACCTCCAATA ATCATGTTAG 2280
 TTTCAGTATA ACATAAAAGC GCTTAAATGA GGATTTAAA AAACGAGTCC GCTTATTTCA 2340
 GACTTCATTT TACTCAGATA TGAATTAGGC ATAAGGTGTC AATTCTGGAT TAATTGGTGT 2400
 ATTAGCTAAG TTGTTGGCAT AGTTACAGAG GATTGCTAGG CTGACACCAA AAACCACATC 2460
 CAAGGCATTT TGTGAGTGT AGCCAGCTTC TAAAACTCA GACAAGGCTT CATCTCCTAC 2520
 ACGACCCTTG GTATTGATAA CTGCCAAGGT AACTTAGCT AGGGTATCCA ATTTAGGATC 2580
 TGTTCATTT GGAGTACGAT TGCGAAGAGC TTGAATCAAG TCATCATFCA TCTGGATTTG 2640
 TTTGATGGAA AAGGCTGTGT GACCTGCGAC ACAGAAGGCA CAACCATTGG TCACGGCTGC 2700
 CGTGATTTGC ACCACTTCAC GCTCAACGGG TGTCAGGCTG TTGCGACGGT GGATAGATGA 2760
 GACAATTTGG TAGGCTTCTA AAACAGTCGG GGCATTGGCC AAGAGACCGA TTAGGTTGGG 2820
 AATATAGCCA TTGTTGTCTT TTTCTACTGT TTCAAGAATT TCTTTCACTT CTGCTGGTGC 2880
 TGACTCTACT GTATGGATAG TAAATGTTGT CATAAGATAC CTCTTTCTT ATTATGACA 2940
 CTAATATTAT TGGAAAATCT TATAAAATCC TGATTCTTAA GTTTATCTAA GATAAAGCTT 3000
 TATTCTCTCA TAAGATTTTC GTTGTATAT TAGTTTATCA CACTTCCAAT CACTTGATA 3060
 ATATATATTA TATATCAGGC TGATAAAAAT TATTTATAGG CAAAAAATC ACACGAGCTG 3120
 TGTGATFCCA TTATTTGTCA AAATACTTTT TAGTTTCAGC AATAACGACT GGCGACAAGA 3180
 CCAAGAGGGC AATCAAGTTT GGCAGAGCCA TCAAGGCGTT AACGATATCT GCGATAATCC 3240
 AGACCATATC CAACTCGATA AATCCTCCTA ACAAGACCAT GAGCACAAA ACCACACGGT 3300
 AGAGCCAGAT AAAGCGAACC CCAAAGAGGA ACTCAAACA GCGTCTCCG TAATAGTTCC 3360
 AACCTAGAAT CGTTGTAAAG GCAAAAAGTA CAAGGAAGAT GGTCAAGAGA GCAGGCCCAA 3420
 AGTGTGAAAA GTTTGTGAG AAAGCTGACT GAGTCAAGGC AACCCCATTC AAGTCACCGC 3480
 TCCAAACTCC AGTTACCAAG ATGGTCAAAC CAGTTAGAGT A 3521

(2) INFORMATION FOR SEQ ID NO: 119:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 1968 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 119:

AACCTGGGCA AGCAAGCTAA AAGCAATGGG ACCTGGAATC CTAATGGCAA CTGCCGCTGT	60
TGGAGGTTCC CACATTGTAT CCTCAACTCA AGCTGGCGGT TCTTACGGTT GGTCTCTACT	120
TCTCTTGGTC ATCTTAGCCA ATGTCTTTAA ATATCCATTT TTCCGTTTGT GTGCTGAATA	180
CACAGCTGAT ACTGGAAAGA CTTTGGTTGA AGGTTATGCC GAAAAAGGAA AACTCTATCT	240
CTGGATTTTC TTTATCCTCA ATGTCTTTTC GGCTATGGTC AACACGGCTG GTGTTGCCAT	300
TCTGTGCTCA GCTATCATCG CCAGTGCCTT CCCAATGATT GGACTTAGCA TTACTCAGTG	360
GTCCCTCATT CTCGTTGCAA TCATTTGGGC TATGCTACTC TTTGGAGGCT ACAAACTTTT	420
AGACGGCATG GTCAAATGGA TTATGTCTGC CTTAACCATT GCGACTGTTC TGGCAGTTAT	480
CATTGCGGCG GTCAAGCATC CAGAATACAG TTCTGATTTT GTCGAGAAGA CACCTTGGCA	540
AATGGCAGCT CTGCCCTTCA TCGTCTCCCT CCTAGGATGG ATGCCGGCTC CTATTGAAAT	600
TTCAGCCATC AATTCACTTT GGTCAGCTGA AAAGAGAAAG ACCGTCAACT TTAACACAGA	660
AGACGCTCTG TTTGACTTTA ACACTGGTTA TATTGGAACA GCTATCCTAG CCGTCTTCTT	720
TGTGGCACTG GGAGCACTGA TTCAGTATCC TACAGGGCAG GCGGTGAAG CTGCTTCAGC	780
CAAATACATC TCTCAATTCG TGGGCATGTA TGCCTCTGTT CTTGGCGAAT GGTCCCGTTA	840
CTTGATTACC TTTATGCGCT TCCTCTGTAT CTTTGGAAACA GTTATAACTG TTATCGATGG	900
CTATCTCGC GTTAATCAGG AATCTCTCCG ACTGCTAATC AGTCAAAAAG AGGACAATCG	960
TAAATCTTTG AACATCTGGA TGACCATCAC TGCTATCATC GGTATCGTCA TTATCAAGTT	1020
CTTCGCTGGT CAGGTTTCAA CCATGCTCCG CTTTGCCATG ATTTGGCTCTT TCCTGACAAC	1080
ACCTTTCTTT GCTCTTTTGA ATTACGCCTT GGTAACGCGT GAAAACAAAA ATCTTCCTTC	1140
TTGGCTCAAA CACCTTGCCA TTGCGGGATT GATTTTCCTC TTTGCTTCGC CATCTTCTTT	1200
ATCTACGCAC TCGCAATCGG AAAAGCAGGG TAAGGGACAA GCGCGAGATG AAGATAAGGT	1260
TTCATTTCAA GAGAAAATTC AGCAAATATT TCTATGATAA AAAGCATAAG AACAGGTTT	1320
TGAAGACCTG AACTTATGCT TTTTACGTT CTAAAGACT GTTTATACTC AAAAAACAGT	1380
TGAACAACCT CAACCACCTC TTATAAGAAC TTTATACTAT TCGAGAATCT CTTCAAACCA	1440

846

CGTCAGCTCT ATCTGCAACC TCAAAGCTGT GCTTTGAGCA ACCTGCGACT AGCTTCCTAG 1500
 TTTGCTCTTT GATTTTCATT GAGTATTAAT TCTCCTTTC CAACTCATAA AAATCTGCCA 1560
 TAATAGCTGC GACATGTTTG ATATCTTCCA GCATGCCTCG CATTTCAAAG TCAGCCAATA 1620
 CAGGGAAGCC AAAGCGTTGA CTGTATTGCT TGGCTGTTAG GCAGTATTGG TTATTAAAGT 1680
 TACGATTTC TGACCCAACC ACACAAAAC ACTTACTAGC ATTGTTACCA TAGGCAATAA 1740
 AATCTCCAC CGGTGTCGTC AAAATCTCAA CATCTCCGTT ATCCACGCCA TTCCACCTT 1800
 CGAGATAGGT CGGCAAAAAA GCGACATAGG GATGGTCCAT TTCATAGAAA TTTTGCCTT 1860
 CCTTGACCAA ATCCTTGATA TGAATCTTTT GAACCTCAAT CCCTTTGTAC TGGGACAAGA 1920
 GATAGTCTTT CAAGCGCGTC ACAAACCTTT CAGTGTGCC ACTCAAGG 1968

(2) INFORMATION FOR SEQ ID NO: 120:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 7172 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 120:

CCGCATTTTT TATCACTAGA CTCGAGACAT CTTTTGAGTG GCTCTTGCTC TCTGGTTTAA 60
 TTTTCTTCCT TGCTCAAGGA CTCCTGCTAT TTCTCTGGT CGTCCGACTC AAACATCAAT 120
 TCGCTGAGAT TTATCCTCAA ATCAATAAAA AGATTGCTT CTAATATTA GGGGTTCTCA 180
 CCATTGATTT TCTATTTTTT GTTCTCTTAG CCTTCATTAG TTCTCAGCGT TTTTCATCTC 240
 TTATGCCAAT CATCACTGCT TGCCATTCTA CTTTTTATTA TATGACAGCT GACTACCTAA 300
 GAGAAAATA TCCAGACTTT TACGACAAAC ACATCTCTTT ATGGGAGTGT CTCTAAAGAA 360
 AAGGAGGTTT TAGCATGAAA AAAATCATCT TCATCAAAAC CATTCAACTC CTGTGCATTG 420
 ATGGAATCAT GCTGGCATTT TTGACATTTA AAAGGGGGCT TACTTGGGAC TGGATTTTGA 480
 TTTATAGCGG TTGGCTCATT TTCTTTCATC CTGTGCTATT GACCTATCTT TCAAACCAAC 540
 TTTGTGACCA CTTTAGTTAA CTCTATTTCC AGATTAGACC GAGATTCTGG CGTTTTGCTT 600
 TACAAATTCT CCTATGGGAT AGCCTGATGA TTCTCTCCTT GGTGTCTTTA AGTGATATTC 660
 CACTTTTCCT TCAGGGAAC CTCCCTATCC TAGGACATCT CATCCCTTCC TATCGCATCT 720
 GCCAAAGCCT GAAAAGAGAC TTCCCCAAG CATATCAAGA ACCGATTTCT TTTTGGAGTA 780
 TTTTATGATA GATGAGAAAG ACCAAGCCGA CTGGGCTTGG TCTTTCCTAT CTCTTTTATG 840
 TATCTAGGAT AATGGTAACA GGTCCATTAT TAACCAGCTC AACCTGCATA TCTGCTCCAA 900

847

AGATGCCTGT CTGAACGGGC ACTTCTTGCG CTAATTTTTG ATTGAAAGCA TCATAGAAGT	960
CTGATGCCAT ATCAGGTTTA GCTGCCCCCTG TAAAGgCTGG ACGATTGCCT CTCTTAGTAT	1020
CCGCAAAGAG GGTAACCTGA GAAATAGAGA GGATTTCTCC TTCAATATCT TTGACAGACA	1080
GGTTCATCTT GCCTTCTGCG TCTGAAAAAA TCCGCATATT GACCAGTTTT CTCACAGCAT	1140
AGTCCAAATC TTCCTCTTGG TCCTCTGGTC CAACACCAAC CAGCAATAAA AGTCCCTGAT	1200
TGATTTTTCC CTGAATCTGG CCTTCTATAC TCACTTGGGC TTTTTTAACC CGTTGGATAA	1260
TGATTTTCAT AATAGCCTTT CTAGTAAGAG CTAGGACAAC TAGCCGTTGG TCCGTTTGAC	1320
AGAGTAAACT TCTGGCACAC TCTTAATTTT ATCGACAACC GTGGTCAGTG TAGAGAGGTT	1380
GGCAATACCG AAGgACACAT GGATATTAGC AACTTCATA TCCTTGGTTG GTTGGGCATT	1440
GACCGTTGAA ATATTCTTGG TTGTATTTGA AAGAACTGC AGTACATCGT TCAACAGTCC	1500
TGTACGGTTG AGACCGTAGA TATCGATATG GGCCATATAC TCCTTATTTG AGCTAGGGTA	1560
CTGGTCTTCC CATTCACAT CAAGGAGACG TTGCTCGTAG TTTTCTTGGG CACGCAGGTT	1620
CATACAGTCC ACACGGTGA TAGCCACACC ACGACCCTG GTAATGTAGC CAACAATATC	1680
GTCACCAGGC ACGGGGTAC AACACTTAGC AATCCGCACT AGGAGACCAG AAGCACCTTC	1740
AATAACCACT CCCCCCTCAT GCTTGACCTT GAGGGTTTCT TTATTTTCAA CCTTGACCTC	1800
GCCACCTTTG ACAAGCTCCT CTGCCTCAGC TTGGCCTTG GCACGCTCTT CCTCACGGCG	1860
TTCTTTTCA GTCAGACGGT TAAAGACGGT AATCGCACCG ATTTCCCCAA AACCAATGGC	1920
CGCAAAGAGG GAGTCTTCTG TCTGTAACT GGTCTTTTGC AGAACTGAT CCATGTGGCG	1980
CTTGTCCATA AATTTATTTG CCACATAGCC ATTTTCTTGG AACTGAGCCA TCAGCATCTC	2040
ACGACCCTTG TTGACAGACA ATTCCTTATC TTGGTTTTTA AAGAACTGGC GAATCTTATT	2100
GCGGCCTTG CTAGTCTTGA CCATATTGAG CCAGTCACGG CTAGGTCCAA AGGAGTTCGG	2160
GTTGGCGATA ATTTCAACCT GATCCCCTGT CTTAACTTG GTTGTCAGTG GAACCATGCG	2220
GCCATTGACC TTGGCACCAG TTGCTTTTTC ACCGACCTTG GTATGGATTT CGTAGGCAA	2280
ATCAATCGGT CCTGAATCTT TGGGAAGGGA ACGGACAGCT CCATCTGGGG TAAAAACGTA	2340
AATCTCCTCA GCCAAATAGT TTTCTTAAC AGAGTCCACA AATTCCTTAG CATCATCAGC	2400
CTGGTCTTGG AGCTCCATCA TCTCCTGAT CCAGTTCATT CCAATAGCTG ATTCCTTGCT	2460
GTAACTTGC CCCTTTATAC CTTTCTTATA AGCCCAGTGA GCCGCAACCC CGTACTCAGC	2520
CACCTCGTGC ATTTCTTGG TTCGAATCTG GAATTCATC GGCCCTTTTG GTCCATAAAC	2580
AGTCGTATGG ATAGACTGAT AACCATTGGC CTGCGGTTG GCGATATAGT CTTTGAAGCG	2640

ACCTGGCATC GGTTCCTAAA ATTCATGCAC GTAACCAAGC ATGGCATAAA CATCACTTTG 2700
GGTATCTAAA ATACAACGAA TAGCAATCAG ATCATAGATT TCCTCAAACC GTTTTCTCTT 2760
GTCCTGCATT TTGCGGAAAA TTGAGTAAAT ATGCTTGGGA CGACCATAAA TCTTCCCTTT 2820
CAAGTGACGT TCTGTCGTAT ACTCCTCTAA TTTTGTGACT ACCTCATCCA CCAAGGCCTC 2880
ACGCTCCCTG CGCTTTTCCCT TCATCATATG GGTAATCTTG TAAACTCCG TTGGATTGAG 2940
ATAACGGAAA GACAAGTCTT CTAATTCCCA TTTGACACTG GAAATCCCA AACGATGGGC 3000
AAGCGGGGCA TAGATTTCCA TGGTTTCTTT GGAAATACGC TCCTGCTTGT CTTTTCGAAG 3060
ATGTTTCAGG GTCCGCATAT TGTGCAAGCG GTCAGACAGT TTGACCAAAA TAACGCGGAT 3120
GTCCTCAGAC ATGGCCATGA GCATCTTGCG ATGATTTTCC GCTAATTGCT CCTCGATCGA 3180
TTTGTACTCG ACCTTGCCAA GCTTGGTAAC TCCGTCAACA ATCATCCGCA CATCAGGACC 3240
AAACTCTCTT TCCAAATCGT CCAAAGTCGC ATCTGTATCT TCCACCACAT CATGCAAGAA 3300
TCCACAAGCT ACTGTTACAG CATCCAGCTT TAGCTTAGCT AAAATACCTG CCACTTGGAT 3360
AGGGTGAATG ATATAAGGCT CGCCTGATTT GCGATATTGA CCACTGTGGC ATTCAACAGC 3420
ATAGACCAAG GCCTTATGGA CAAAATGAAC ATCCTCTTCC GTTAAATATF CTTTGGTTAA 3480
AGCGACAACF TCTTCGCCTG TTAAATTCAC TTCTTTCGGC ATCTCTACTC TCCAATTCTT 3540
CCTACCATTT TATCACTTTT TTAAGAATAT GAAAACCTAGA TTGGAACAGA ATAAGAAAAA 3600
AATAATTCAA AATGCTTGA TAATCTGAA TTATTGGTCC GTAATATACT ACGAAGTTAG 3660
ATTTTAAACT TAGGTGATAG AAGGAGAGAT AGAAGAACGG AAACCATATF GTAACCCAAA 3720
GACTTTCTGA CTTCCTCAAT TCCATTGAAG ATACGAAAGA TAAACGGTGG AACTCGTATC 3780
ACATACACTG GTACCTTGAC TGGATTTTGG AATTAATACT AAATGAAAAT CAAAGAGCAA 3840
ACTAGGAAAC TAGCCGCAGG TTAATCAAAG CACCGCTTTG AGGTTGCAGA TAAAGTTGAC 3900
GCGGTTTGAA GAGATTTTGG AAGAGTATAA AAATCCTCAA GATACTTTCT TCTATCCTTT 3960
AGTTTATAAG GAGAATACCT ATGAAAAAAA CTGCTATTTT TATCTTTGCT CTCCTAATGT 4020
TAGGAGTTTG CTGCTGTTC CTATTCAGCC AGCAAAGCTA TAAAAACAG TCGTTCAATA 4080
CTATGCTAAC GACCAGAACC TGCCAGTAG GATAACTTAT AGTGAATATA GCGACAAATG 4140
AGAAGCCAAC TACGGTAGCA CTCTAAACAT CACGTCTATC AAACAAGCTA ATGACGGAGT 4200
TTATGCAACC TATGAAGGGC AATTGACACC TTCCAATAT TGATAAATG ATAACCAGCC 4260
TGTCTTCATC TAGTCATGCT GGTTTTTAAG TTCATTTTAA ATCCTTACCT ATTCTCCCTA 4320
ACTGTGCTAT ACTTAATTTA TACTCAATGA AAATCAAAGA GCAAACCTAGA AAGCTAGCCG 4380
CAGGCTGTTT AAAGCACTGC TTTGAGGTTG CAGATAAAGT TGACGCGGTT TGAAGAGATT 4440

TTCGAAGAGT	ATTAGTACAT	TCTTTGAGAT	TGGAGCTAGT	ATGAAAATCC	ATAAAACCGT	4500
GAATCCTGTT	GCCTATGAAA	ATACCTATTA	TCTAGAAGGC	GAAAAGCACC	TCATCGTCTG	4560
CGATCCTGGT	AGTCATTGGG	AAGCCATTCC	TCAGACAATC	GAGAAGATCA	ACAAACCGAT	4620
CTGTGCTATT	CTCTTGACCC	ACGCCCATTA	TGACCATATC	ATGAGTCTGG	ACTTGGTTCG	4680
CGAGACGTTT	GGCAATCCTC	CTGTCTATAT	CGCAGAGAGC	GAAGCCAGCT	GGCTCTACAC	4740
TCCTGTTCGAT	AATCTCTCCG	GTCTCCCTCG	CCACGATGAT	ATGGCAGATG	TGGTCACAAA	4800
ACCTGCAGAA	CACACCTTTG	TCTTTCACGA	AGAATACCAA	CTAGAGGAAT	TTCGTTTTAA	4860
GGTCTTACCG	ACCCAGGGC	ACTCTATCGG	TGGTGTTC	CTAGTCTTTC	CTGATGCTCA	4920
TCTAGTCTTG	ACGGGAGATG	CTCTATTCCG	CGAACTATC	GGACGGACCG	ACCTTCCGAC	4980
TGGTAGCATG	GAGCAACTCC	TTCATAGTAT	CCAGACCCAA	CTCTTCACCC	TACCAAATA	5040
CGATGTCTAT	CCAGGACATG	GTCCAGCTAC	TACTATCGCT	CACGAAAAGG	CCTTCAATCC	5100
CTTTTCTAG	CAAGATGATG	ACAATCGAAA	TTAAGTAAA	CTATCCAGCA	AATCTTTCTA	5160
TTACAAAAGG	CATCCTATCA	AGGTTTTCAC	ACATGATTGG	ATGCCTTTTT	TCTGATGACT	5220
AGATTTTTTG	CATTACCAA	TAATCACGCG	CTCCTCTGGT	GAACGCCACA	TCCGTCTCC	5280
TTCTTTGACA	TCATAGGTTG	TAAAGAAATC	GTGGAAGTTT	GGTACTTGCA	CATTGACACG	5340
GAGTTTGGCT	GGTGCCTGCA	CATCGACGCT	AGCCAAAAGT	TTCATAAAT	CTGGTCGACC	5400
TTTCATGCGC	CAGATGCGAC	CGAAGTTGTA	GAAGAACTCT	TCTGCTGAGA	AGTCTGCTTC	5460
TCTCTTAGCT	GCTTCAAGCG	CTGCTGCGAT	TCCTCCCAAG	TCAGCCACGT	TTTCTGATAC	5520
AGTCAATTTA	CCGTTAATGG	TTGCTCCATA	AGAATCCTGT	CCATCAAAT	GGTCAATGAC	5580
TTTTTGTGTT	TTCTCCTTGA	AGGCAGCATA	GTGCTCTCT	GTCCACCAAT	CCTTGAGGCT	5640
ACCATTTTTCG	TCAAAGGAAG	CCCCGTTAGT	ATCAAAGGCG	TGGGAAATTT	CATGGGCAAT	5700
CACTGCCCCA	ATACCACCGT	AGTTAGCAGA	AGATGACTGA	TGCAAGTCAT	AGAAAGGCGC	5760
CTGTAAAATG	GCCGCTGGAA	AGACAATCAG	GTCTTCTGA	GGATTGTAGT	AGGCATTGAC	5820
CATATGAGCA	GGCATGCCCC	ATTCCTTATA	ATCTACAGGC	TGGTCCACT	TACTCCAAT	5880
GTGCTTGATT	TCCACACGCG	CAAAGGCTAG	AGCATTCTCA	AAAAGACTGG	CAGTTTCATT	5940
CACTACCTTA	TCCTTGTAAC	GTGCAGGCAA	TTCTTCTGGA	TAGCCAATAT	AAGGTTTGAT	6000
CACATTGAGC	TTACAGATAG	CCTGTTTACA	GGTTTCTGGA	GTGAGCCAGT	CATTCTTAAG	6060
CAGACGCTCC	TTATAAACAT	CAATCATGGT	TGCCACTTTT	TTCTCCACAT	CCGCCTTGGC	6120
TTCTGGAGAG	AACTTCTCAC	GGGCGTACCA	AAGACCCAGG	GCTTGCTTGA	AAGGTTCTTG	6180

850

TGCTAGATGA TAAGCTGCTT TGACCTTATC TTTTGCCTCT GGAACTCCAG AAAGGGCACG 6240
GCTGTAGGCA CCAGACAAA CACGGATATC CTCTGTTAAA TAGCTGGTTG AAAGATTGAC 6300
AAACTCAAA ATCAAGGTTG CTTTAAGGAG AGACCAGGCT TCCTCACTGT AGAATTGCTC 6360
TGCTGCTTGC CAGAAACGTT CCTCGTCTAC AATAACCTTG TCTGGTAATT GCCCAATAAC 6420
TGCTTTGAAG AAGTCATCCA AAGGTAGGGC AGGCGCGAAT TTCTTGAAAT CTTCGTAAGA 6480
ATATGGATGA TAGAGTTTAG CATATTCTGA ACTTCTTCA TTAGAGAGCA CCACTGCCGC 6540
AACTCGGCGG TCCAATTCAA GTCTTTTTTC TAGCAAGTCT TCAATTTCTT CATCAGAGAA 6600
ATCATAAGCC TTGAGGAGAT TTGCGCTGCT TTCTTTCCAA AGAGTCAAGA GCTCTTCGCG 6660
CTGAGGATGT TCTTCTGCAT AGTAGGTCGT ATCTGGCAAG ATTGTGCTTG GAGCGCTAGC 6720
CCATAGAACA TTGATTCTAG CATCCATAAA GTCTGGCGAT ACACCAAAG GAAGGAAGTT 6780
TGGTTTTCTT GCAAGCTCAA ACTCTGCTAG TTTAGCTGTA AAATCCGCAA AAGTCTCCAA 6840
TTCTTGGAAT TCTTTAAGGA GTGGTAAGAC AGGTGTGATA CCGTCAGCTT CTCTCTTGTC 6900
AAAATCACGA ACTAGGCGGT GGTATTTGAC AAAGTTTTCC AAGATAGCAT CCTCAGGCAC 6960
TTCTTCACCT GCTAACCACT TGTCTGTGTG CGCCAGCATC AGGTCTTCAA TTTCTCTGTC 7020
TAAATCAACA AAACCTCTG TTTGAGACTT ATCTGCTGGG ATTTAGCTG TCTGTTGCCA 7080
TTCTCCATTG ATAGCATCAT AAAAATCATC TTGATAACGT GTCATCTTGT TCTCGCTTTC 7140
ATTTGTATTT GCATTTATCT TAACAAAAT CG 7172

(2) INFORMATION FOR SEQ ID NO: 121:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 4518 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 121:

CGGGAAGTTA TGCGATCTAG ACTTCGTTCC TGTACAGCTA CTTTCTCAGG TGGTCTTGTT 60
GTTTGTATGA GTTTGTTTAG AGAGGATCTT TCTATGTCTT TCTTCTTAT TTTTGTTTTA 120
TATGCTTTTC TGATTTCTTA TCTAATTTAT GGTATTTC A GACTAAAAG GAAATACCGA 180
GTAGATGAAT AGCAAGGTTT TAGGTCTTCA GATTGATTTT TAGCACTCTT GATAAAAGAG 240
TGCTAATTTT TTGAGTTTTT GTCTTGACAT TCTCTTCTAA GGGTGTATAA TAGAATCATG 300
AGTTAGCACT TGGATGCATT GAGTGCTAAT TGATCAGACA GAGAGGAGTG ATGAGATGGT 360
TACAGAGCGT CAGCAGGATA TTTTAAATCT GATTATTGAC ATCTTTACCA AAACGCACGA 420

ACCTGTCGGA	TCAAAAGCCT	TGCAAGAGTC	TATTAACTCT	AGCAGTGCAA	CCATTCGTAA	480
TGACATGGCG	GAAC TAGAAA	AACAAGGTT	GCTTGAGAAG	GCTCATACTT	CAAGTGGTCG	540
GATGCCAAGT	GTTGCTGGTT	TTCAGTACTA	TGTGAAACAC	TCACTGGATT	TTGACCGGCT	600
GGCTGAAAAAT	GAGGTATATG	AGATTGTCAA	AGCCTTTGAT	CAGGAATTCT	TCAAATGGGA	660
GGATATTCTG	CAAGAGGCTG	CTAACTTACT	AACAGACCTG	AGTGGCTGTA	CGGTAGTGGC	720
ACTGGATGTT	GAGCCGAGCA	GGCAACGTTT	GACAGCCTTT	GATATCGTTG	TTTTGGGGCA	780
ACATACAGCC	TTGGCGGTAT	TTACCCTAGA	CGAGTCGCGA	ACGGTFACTA	GTCAGTTTCT	840
GATCCAAGG	AACTTCTTGC	AGGAGGATTT	GCTGAAACTG	AAGAGCATCA	TTCAGGAACG	900
TTTCCTCGGT	CACACCGTTT	TAGATATTCA	CTACAAGATT	CGGACGGAGA	TTCCGCAGAT	960
TATCCAGCGT	TACTTTACAA	CAACGGATAA	TGTCATCGAT	CTCTTTGAAC	ACATCTTTAA	1020
GGAAATGTTT	AACGAAAACA	TTGTGATGGC	GGGCAAGGTC	CATCTCTTGA	ATTTTGCCAA	1080
TCTAGCAGCC	TATCAGTTCT	TTGACCAACC	GCAAAGGTG	GCCTTGGAGA	TTCGTGAGGG	1140
GTTGCGTGAG	GATCAGATGC	AAAATGTTTCG	TGTTGCAGAC	GGTCAAGAGT	CCTGTTTAGC	1200
TGACCTAGCG	GTAATCAGTA	GTAAGTTCCCT	CATTCCTTAT	CGGGGAGTTG	GAATCTTAGC	1260
CATTATCGGT	CCAGTTAATC	TGGATTACCA	ACAGCTAATC	AATCAAGTCA	ATGTGGTCAA	1320
CCGTGTTTTG	ACCATGAAGT	TGACAGATTT	TTACCGCTAC	CTCAGCAGTA	ATCATTACGA	1380
AGTACATTAA	GATTGAAATC	ATTAAAGGAG	GCGAACATGG	CCCAAGATAT	AAAAAATGAA	1440
GAAGTAGAAG	AAGTTCAAGA	AGAGGAAGTT	GTGAAAACAG	CTGAAGAAAC	AACTCCTGAA	1500
AAGTCTGAGT	TGGACTTGGC	AAATGAACGT	GCAGATGAGT	TCGAAAACAA	ATATCTTCGC	1560
GCTCATGCAG	AAATGCAAAA	TATCCAACGC	CGTGCCAATG	AAGAACGTCA	AAACTTGCAA	1620
CGTTATCGTA	GCCAGGACTT	GGCAAAAGCA	ATCTTACCAT	CTCTTGACAA	CCTTGAGCGT	1680
GCACTTGCAG	TTGAAGGTTT	GACAGATGAT	GTGAAGAAGG	GCTTGGGGAT	GGTGCAAGAA	1740
AGCTTGATTC	ACGCTTTGAA	AGAAGAAGGA	ATTGAAGAAA	TCGCAGCAGA	TGGCGAATTT	1800
GACCATAACT	ACCATATGGC	CATCCAAACT	CTCCCAGCAG	ACGATGAACA	CCCAGTAGAT	1860
ACCATCGCTC	AAGTCTTTCA	AAAAGGCTAC	AAACTCCATG	ACCGCATCCT	ACGCCCAGCA	1920
ATGGTAGTGG	TGTATAACTA	AGATATAAAG	CCCGTAAAAA	GCTCGCAGTA	AAAATAGGAG	1980
ATTGACGAAG	TGTTTCGATGA	ACACAAGAAA	ATCTATCTTT	TTTACTCAGA	GCTTAGGGCG	2040
TGTTTCGATTC	GGCAATTCTG	ACGGTAGCTA	AAGCAACTCG	TCAGAAAACG	GCAATCGCTA	2100
TGGCGTTTGC	CTAGCTTCCT	TACTAACTCG	TCGTCGAAAT	AAAATCGATT	TCGACTCCTC	2160

GTGTCGCAAT	TTACATAATA	GAAAACCTGT	CCGAAACGAC	AATAAACTAT	GAAGAAAGAT	2220
AAAATATGTT	TGGCTTTGTA	ATAGTGAGCG	AAGCGAACCA	AACACGATAC	TCTTCGCCGT	2280
GGCGCTATTT	GCGCAAATTT	TGAGACCTTA	GGCTCAAAGT	TTAGTCAAAG	AGATTGACGA	2340
AGTCAAGCTC	TGACGGCGTC	GCCACTGTCT	CCACTTAAGA	AGAGTATCAA	AAAGAAAAAT	2400
AGAAAAFTAA	CTAACAAAGGA	GAAAAACACA	TGTCTAAAAT	TATCGGTATT	GACTTAGGTA	2460
CAACAAACTC	AGCAGTTGCA	GTTCTTGAAG	GAACTGAAAG	CAAAATCATC	GCAAACCCAG	2520
AAGGAAACCG	CACAACTCCA	TCTGTAGTCT	CATTCAAAAA	CGGAGAAATC	ATCGTTGGTG	2580
ATGCTGCAAA	ACGTCAAGCA	GTTACAAACC	CAGATACAGT	TATCTCTATC	AAATCTAAGA	2640
TGGGAACTTC	TGAAAAAGTT	TCTGCAAATG	GAAAAGAATA	CACTCCACAA	GAAATCTCAG	2700
CTATGATCCT	TCAATACTTG	AAAGGCTACG	CTGAAGACTA	CCTTGGTGAG	AAAGTAACCA	2760
AAGCTGTAT	CACAGTTC	GCTTACTTCA	ACGACGCTCA	ACGTCAAGCA	ACAAAAGACG	2820
CTGGTAAAT	TGCTGGTCTT	GAAGTAGAAC	GTATTGTTAA	CGAACCAACT	GCAGCAGCTC	2880
TTGCTTATGG	TTTGACAAG	ACTGACAAAG	AAGAAAAAAT	CTTGGTATTT	GACCTTGGTG	2940
GTGGTACATT	CGACGTCTCT	ATCCTTGAAT	TGGGTGACGG	TGTCTTCGAC	GTATTGTCAA	3000
CTGCAGGGGA	CAACAAACTT	GGTGGTGACG	ACTTTGACCA	AAAAATCATT	GACCACTTGG	3060
TAGCAGAATT	CAAGAAAGAA	AACGGTATCG	ACTTGCTTAC	TGACAAGATG	GCAATGCAAC	3120
GTTTGAAAGA	TGCGGCTGAA	AAAGCGAAGA	AAGACCTTTC	TGGTGTAACT	TCAACACAAA	3180
TCAGCTTGCC	ATTTATCACT	GCAGGTGAGG	CTGGACCTCT	TCACTTGGA	ATGACTTTGA	3240
CTCGTGCGAA	ATTTGACGAT	TTGACTCGTG	ACCTTGTTGA	ACGTACAAAA	GTTCCAGTTC	3300
GTCAAGCCCT	TTCAGATGCA	GGTTTGAGCT	TGTCAGAAAT	CGACGAAGTT	ATCCTTGTTG	3360
GTGGTCAAC	TCGTATCCCT	GCCGTTGTTG	AAGCTGTTAA	AGCTGAAACT	GGTAAAGAAC	3420
CAAACAAATC	AGTAAACCCT	GATGAAGTAG	TTGCTATGGG	TGCGGCTATC	CAAGGTGGTG	3480
TGATTACTGG	TGATGTCAAG	GACGTTGTCC	TTCTTGATGT	AACGCCATTG	TCACTTGGTA	3540
TCGAAACAAT	GGGTGGAGTA	TTTACAAAAC	TTATCGATCG	CAACACTACA	ATCCCAACAT	3600
CTAAATCACA	AGTCTTCTCA	ACAGCAGCAG	ACAACCAACC	AGCCGTTGAT	ATCCACGTTT	3660
TTCAAGGTGA	ACGCCCAATG	GCAGCAGATA	ACAAGACTCT	TGGACGCTTC	CAATTGACTG	3720
ATATCCCAGC	TGCACCTCGT	GGAATTCCTC	AAATCGAAGT	AACATTTGAC	ATCGACAAGA	3780
ACGGTATCGT	GTCTGTTAAG	GCCAAAGACC	TTGGAECTCA	AAAAGAACAA	ACTATTGTCA	3840
TCCAATCGAA	CTCAGGTTTG	ACTGACGAAG	AAATCGACCG	CATGATGAAA	GATGCAGAAG	3900
CAAACGCTGA	AGCCGATAAG	AAACGTAAAG	AAGAAGTAGA	CCTTCGTAAT	GAAGTAGACC	3960

853

AAGCAATCTT TGC GACTGAA AAGACAATCA AGGAAACTGA AGGTAAAGGC TTCGACGCAG 4020
 AACGTGACGC TGCCCAAGCT GCCCTTGATG ACCTTAAGAA AGCTCAAGAA GACAACAAC 4080
 TGGACGACAT GAAAACAAAA CTTGAAGCAT TGAACGAAAA AGCTCAAGGA CTTGCTGTTA 4140
 AACTCTACGA ACAAGCCGCA GCAGCGCAAC AAGCTCAAGA AGGAGCAGAA GGCGCACAAG 4200
 CAACAGGGAA CGCAGGCGAT GACGTCGTAG ACGGAGAGTT TACGGAAAAG TAAGATGAGT 4260
 GTATTGGATG AAGAGTATCT AAAAAATACA CGAAAAGTTT ATAATGATTT TTGTAATCAA 4320
 GCTGATAACT ATAGAACATC AAAAGATTTT ATTGATAATA TTCCAATAGA ATATTTAGCT 4380
 AGATATAGAG AATTATATTA GCTGAACATG ATAGTTGTAT CAAAAATGAT GAAGCGGTAA 4440
 GGAATTTTGT TACCTCAGTA TTGTTGTCTG CATTTGTATC GCGGATGGTA CCGTATCTGA 4500
 CGAACGTTCA GCTTATAT 4518

(2) INFORMATION FOR SEQ ID NO: 122:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 8145 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 122:

TGCTATTTTC GATTCCCTTG GCGTTTTGA TTGCCTTTGC CTTGCAAGTC CATTGAAGC 60
 CCCTCCATTA TCTGATTAAC ATTTACATCT GGGTTATGCG AGGAACCCCC TTA CTCTTG C 120
 AACTGATTTT TATCTATTAT GTGCTCCCAA GTATTGGGAT TCGTTTAGAC CGCCTTCCTG 180
 CAGCTATTAT TGCTTTTGT CTCAACTATG CAGCTTACTT TGCAGAAAT TCCGTGGGG 240
 GAATGACAC TATPCCAAGA GGACAGTATG AGGCCGCCAA GGTCTTGAAG TTTAGCCCTT 300
 TTGACAGAGT GCGCTATATT ATCTTGCCCC AAGTGACCAA GATCGTTCTT CCTAGTGTCT 360
 TTAATGAAGT TATGAGTTTG GTCAAGGATA CTTCTTTGGT CTATGCTCTC GGAATTT CAG 420
 ACCTTATCTT GGCTAGTCGA ACAGCTGCTA ACCGCGATGC TAGTCTAGTT CCTATGTTCT 480
 TGGCAGGAGC CATTTATTTG ATTTGATTG GGATTGTGAC AATTATTTCC AAAAAAGTTG 540
 AGAAGAAGTA TAGTTATTAT AGATAGGAGG CTGCCATGTT AGAATTACGA AATATCAATA 600
 AAGTCTTTGG AGACAAACAA ATCCTGTCTA ATTCAGTCT AAGTATTCCT GAAAAGCAAA 660
 TCCTGGCTAT CGTTGGACCT TCTGGTGGAG GTAAGACAAC TCTTTTACGT ATGCTTG CAG 720
 GTCTTGAAAC CATTGATCA GGGCAAATCT TTTATAATGG ACAACCTTFA GAGCTGGATG 780

854

AATTGCAGAA	GCGCAATCTA	CTGGGATTTG	TCTTCCAAGA	TTTTCAACTA	TTTCCTCATC	840
TATCAGTTCT	GGAAAATTTG	ACTTTATCGC	CTGTGAAGAC	CATGGGAATG	AAGCAGGAAG	900
AGGCTGAGAA	GAAGGCGAGT	GGACTCTTGG	AACAGTTAGG	ACTAGGAGGA	CACGCAGAGG	960
CCTATCCTTT	CTCACTATCT	GGTGGGCAAA	AGCAGCGGGT	GGCTTTGGCG	CGTGCTATGA	1020
TGATTGACCC	AGAAATCATT	GGCTACGATG	AACCAACTTC	TGCCCTGGAT	CCAGAATTAC	1080
GTTTGGAAGT	GGAGAAGCTA	ATCTTGCAAA	ATAGGGAACT	TGGGATGACC	CAGATTGTGG	1140
TTACCCATGA	TTTGCAGTTT	GCTGAAAATA	TCGCAGATGT	ATTATTGAAA	GTAGAACCTA	1200
AATAGGAGGA	AAAATGGATG	AAAAAATGGA	TGCTTGATATT	AGTCAGTCTG	ATGACTGCTT	1260
TGTTCTTAGT	AGCTTGTGGG	AAAAATCTA	GCGAAACTAG	TGGAGATAAT	TGGTCAAAGT	1320
ACCAGTCTAA	CAAGTCTATT	ACTATTGGAT	TTGATAGTAC	TTTTGTCCA	ATGGGATTTG	1380
CTCAGAAAGA	TGGTTCTTAT	GCAGGATTTG	ATATTGATTT	AGCTACAGCT	GTTTTTGAAA	1440
AATACGGAAT	CACGGTAAAT	TGGCAACCGA	TTGATTGGGA	TTTGAAAGAA	GCTGAATTGA	1500
CAAAGGAAC	GATTGATCTG	ATTTGGAATG	GCTATCCGC	TACAGACGAA	CGCCGTGAAA	1560
AGGTGGCTTT	CAGTAACTCA	TATATGAAGA	ATGAGCAGGT	ATTGGTTACG	AAGAAATCAT	1620
CTGGTATCAC	GACTGCAAAG	GATATGACTG	GAAAGACATT	AGGAGCTCAA	GCTGGTTCAT	1680
CTGGTTATGC	GGACTTTGAA	GCAAATCCAG	AAATTTTGAA	GAATATTGTC	GCTAATAAGG	1740
AAGCGAATCA	ATACCAAACC	TTTAATGAAG	CCTTGATTGA	TTTGAAAAAC	GATCGAATTG	1800
ATGGTCTATT	GATTGACCGT	GTCTATGCAA	ACTATTATTT	AGAAGCAGAA	GGTGTTTTAA	1860
ACGATTATAA	TGTCTTTACA	GTTGGACTAG	AAACAGAAGC	TTTTGCGGTT	GGAGCCCCTA	1920
AGGAAGATAC	AAACTTGGTT	AAGAAGATAA	ATGAAGCTTT	TTCTAGTCTT	TACAAGGACG	1980
GCAAGTTCCA	AGAAATCAGC	CAAAAATGGT	TTGGAGAAGA	TGTAGCAACC	AAAGAAGTAA	2040
AAGAAGGACA	GTAAGATAAA	ATAGTGGCTG	AAACTGCGTT	TTGATTAGCA	AAACGTAGTT	2100
TTTTTTGTAA	TCTAGGAAAA	CGATAATAGC	GATTGAATAT	GGATAATTGA	ATATGGAATA	2160
GCCCACTGTG	ATTTCTAAAA	CATTGTAAAA	AATTGATTTG	ACTTCCAAAA	TTAAAATGTT	2220
CTGTAATGAA	ATACTGATGT	AACTGTTTTA	GGAACAATAA	AACGCATAAT	ATCAAGGTTT	2280
TTGCACCTTA	CATTATGCGT	TTTTGTGATT	TTAAGACTTG	TTAGCTGATT	TTTTACAATC	2340
CTGCGAAATC	TTTGATTTCT	TGTGCTGACA	TTGAAGAGTC	GCAACGGACG	TTGATTGTGC	2400
CATCTGTAAT	ATGAACAAAA	CCTGGTACAG	TTGGGATTCC	ATAGCGTGAG	CGGAATGCTT	2460
GCAAATCATF	GAGTTGGCTT	GGTCTTCAC	TATTGATGAA	GTAATGTGA	GCTTTGGTTT	2520
CAGTACGAC	ACCTGACAA	GTACCTGCAA	ATTTACGGCA	GTAAGGGCAA	GTTTTGCGAC	2580

855

CGATAAAGAA	GGTTGCAGTT	TCTTTTTTAT	CAAGAGCTTC	TTGCGCACGC	ACAACTGTAG	2640
TGACTTCAAG	GTCTTTGATG	TTATCTAAAA	ATTGTTCCAT	GAGATTACCT	CGCTTTCATT	2700
GATAAGTCTA	GTATGCCATA	AAGTTTCTAA	AATTGCTTAG	ATTTGATACG	AAAAAAGATG	2760
AGGTTGGTTG	GTCTCATCTT	TTATAGGTCT	TTATTTTACA	AATGCATTGA	TTTCTGCTTC	2820
GATGTTAGCA	ATCTTAGCTT	GTGATCTTTC	GTTGGTTTCC	CCTACAACCTG	CAATGTAGAA	2880
CTTGATTTTT	GGTCTGTAC	CTGAAGGGCG	AACGGCAATC	CATGAACCGT	CAGCAAGTGT	2940
GTATTTCAAC	ACATCACTTG	GAGGAGTTGT	CAAGTTTGTA	ACAGTACCGT	CAGCAACAGT	3000
AGCAGTTTGT	GCCTTGAAGT	CTTCTACGAC	AGTGATAGCT	GTTGCGTTCC	ATTCTGTTGG	3060
AGCATTGTTG	CGGAATTTAG	CCATAATCGC	TTTGATTTGT	TCAGCACCAT	CGACACCTGA	3120
AAGAGTAACA	GAGATTGTTT	TTTCTGCGTA	GTAGCCATAT	TCTTTATAGA	TTTCTTCGAT	3180
ACCGTCAGCA	AGTGTCAAAC	CACGAGAACG	GTAGTAGGCA	GCAAGTTCAG	CAACTACAAG	3240
AACGGCTTGG	ATGGCATCTT	TATCACGTAC	AAATGGTTTA	ATCAAGTAAC	CGAAGCTTTC	3300
TTCAAATCCC	ATCATGTAAG	TGTGGTTGTG	TTTTTCTTCG	AATCTCTGGA	TTTTTTCAGC	3360
GATAAATTTG	AAACCTGTCA	AGACGTTGAA	CATAGTTGCG	CCGTAGCTTT	CAGCAATCTT	3420
CGTTACCAAG	TCAGTTGAAA	CGATAGATTT	GCAGAGAGCG	GCATTTTCAG	GAAGAGTTCC	3480
AGCGTTTTTG	TGAGCTTCCA	AGATGTATTT	AGCCATGATA	GCACCGATTT	GGTTACCTGA	3540
AAGGTTGAGG	TAGCTACCAT	CTTTTTGAAG	AACTTCAACA	CCAACACGGT	CAGCGTCTGG	3600
GTCAGTTGCG	ACAAGAACAT	CTGCACCAAC	TTGACGACCA	AGTTCTTCAG	CAAGGGCAAA	3660
GGCTGCTTGG	CTTCTGGGT	TTGGAGATGT	TACAGTTGAA	AAGTCTGGGT	CAGCAGTTGC	3720
TTGCGCTTCA	ACAACTTGAA	CAGAGTCAA	TCCTGCTTGG	GCAAGAGCAC	GACGAGCCAA	3780
CATTTACCA	GTACCATGAA	GTGGTGTGTA	GACAATCTTC	ATGTCTTTAC	CAAATTCCTC	3840
AATCAAGGCT	GGGTTGATGT	TTATGTCCCT	AACCTCTTTA	AGGTATTCTA	TGTCAACAGC	3900
TTCCGCCGATA	ACTTCAATCA	AGCCAGAAGC	TTTTTCAGTT	TCCACATCAG	CAACTTCAAC	3960
TGCAAATGGG	TTTTCGATTG	CACGGATATA	AGTAGTCAA	GCGTCCGCAT	CGTGTGGAGG	4020
CATTTGTCCA	CCGTCTTCAC	CGTAAACCTT	GTAACCGTTA	AATGGAGCAG	GGTTGTGGCT	4080
GGCTGTGACC	ATGATACCTG	CGAAACAGTT	GAGATGACGA	ACTGCAAATG	ATAGTTCTGG	4140
AGTCGGACGA	AGGCTTTCAA	ATACGTAAGA	TTTGATGCCG	TGTTTAGCAA	GAACTGCCGC	4200
AGATTCAAAG	GCAAACCTCAG	GTGAGAAGTG	ACGGCTATCG	TAGGCAATTG	CTACACCCGC	4260
TTCTTTCTCG	TTTCCACCTT	TTGACTCAAT	CAAACGAGCC	AATCCTTCAG	TAGCTTGGCC	4320

AACAACGTAG	ATGTTGATAC	GGTTTGTACC	AGCACCAACC	AAGCCACGCA	TACCTGCAGT	4380
ACCAAATTCA	AGATTTGTAT	AGAAGGCATC	TTCTTAGTT	TTTTCGTCCA	TATTTTCCAA	4440
ATCTTGACGA	AGGTAGTCAC	GAAGCTCCAC	AAAATCAACC	CATTTCTGGT	AATTTTCTTG	4500
GTAAGACATT	CAAATCTCC	TTTATTTTAA	AAACATTTAA	TCAGTTAAT	TATATCATTT	4560
TTTTTAGTTT	TAGTAAAACC	TTATCTGCTT	CGAACATCTC	TTCAAACCAG	GTCAGATTGA	4620
ATTTTGGGGT	TATATGATGT	TGAGGCTAGG	AAAAATCAA	TTTCAGTAAA	AAAAGTAAGT	4680
CTTCTCATAA	CAAACATTG	ATATAGTTAC	TTAGTTTAA	ACAAGCATAT	TATAATAAAG	4740
CTATGGCATA	TAGTACTGAT	TTTAAACAGC	GAGCATTAGA	TTACATCAAA	GAGGGGCACA	4800
GCCATGTCGA	GGCAGCCAAG	TTTTTTGGTG	TTGGCGTCAG	AACTCTCTTC	ACGTGGGAAA	4860
AGAAAGACGT	GAACAAGAAC	ACATAGAGAG	GAAAAAGCGA	GTCGTCAAAA	ACCGAAAGAT	4920
TCCTTTAGAG	GAATTGAAAG	CCTTTGTAGA	GGCTCATCCA	GATGCTTTTT	TACGGGAAAT	4980
TGCGGCACAT	TTTGATTGTG	CTGTTCTTTC	AGTATGGGCA	GCTTTAAAGC	AGATTAAGGT	5040
CACTTTAAAA	AAAGATGACG	AGCTTTAAGG	AACAAGACCC	AGAAAAGTAG	CCTTATTTCT	5100
TAAGAATTTT	AATAGTTTAA	AGCACCTAGC	ACCTGTTTAT	ATTGATGAAA	CAGGAATCGA	5160
CCGCTATCTC	TATCGTCCTT	ATGCAGGGGC	TCCTAGAGGG	GAGAAAGTCT	ATGAAAAGAT	5220
TAGCGGACGT	CGTTTTGAGC	GAAC TTCAAT	TGTTGCAGGA	CAAGTAGACG	GAGAGTTTAT	5280
AGCTCCCATG	ATTTACAAGA	AAAGCATGAC	AAGCGATTTT	TTTGTGGAGT	GGTTCAAAAC	5340
GCAACTCCTA	CCTGCTTTGA	AGACACCTCA	TGTTATTGTC	ATGGGCAATG	CTGGTTTTC	5400
TCCCAAGAAC	ATTTTGGATG	AACTCTGCAT	CCAAGATAAA	CACTTTTCT	TACCTCTACC	5460
ACCTTATTCA	CCGGATTTGA	ATCCTATTGA	GCAAGCTTGG	GCTATCTTGA	AAAAGAAAGT	5520
GACGGATGTA	TTAAGGGAAG	TTCCAACAT	TTTTGAATGT	TTGGAATGCT	TTTTTAAAC	5580
TAGATGACTA	TAACGGTTCT	AAAGGAACCT	ATCGAGTAGT	CATTAAAACT	AAGGATACTG	5640
CTGGTTAAGA	GAAGACGGTA	TACAATCAAA	CCATTCACCG	TGTAGCCGAA	ATCGTTCAGA	5700
ATGAAGACTT	GTATCAGAAAT	GAAGACTTGT	ATAAGAAAGG	TTTGAATGTT	GAAC TTGCGC	5760
ACCAACAAAT	TAAGGGATTT	TTTGAAGCAG	AGTTTAAAAA	TCGTATTAAT	GGAGTTCTTA	5820
ATACTAAAAAT	AAAAAATAGT	ACATTAATC	GTGTAAATAA	AAAAACTATA	CACCAGAGCA	5880
ACAAAAACTC	CATGATCAAT	TTGAAGCAGA	AGCAACGGAA	GATGCTAAAA	AACAAGCGCA	5940
TATTGTGTTG	AATGTTGACC	AGGATTTTAT	GAGCATATCT	AAGTCTAATA	AAAGTGGTTC	6000
AGACTGGAAG	AAAAC TTCA	CAGTGAGGAT	AACCAATAGG	CTAGCAAATG	ACTTGAATAA	6060
TGTCTTGAAA	CAGGTTGATA	AAGATACTCC	TAATACCCCA	ACTTGGCTAA	ACTCAGCTGC	6120

TTCTAAAGCT	AAAGATGATG	ACAGAGTATA	TAAACTACTG	AAGACTCTTA	TACCAGGAGA	6180
AAATTACCTA	TCATGTTAAG	GATAATCAGC	TAGAAGTAGA	AACAGATAAA	TACACATATA	6240
CTGCCGCTAG	AAATGGTAGT	AAGGAAGTGT	GTATTCAAGA	GTCAGATATA	GCAGCAACTC	6300
TAAGTGCCGA	TGAATATAAT	TCTAATCGCC	AAACTTTTGA	GAGAGAATAC	AAATACAAAA	6360
GCAAATGCC	TTAATAATGG	TTGGGCTAGA	TCTGGTCTG	AAGAGTCAA	AAAGTTCTCC	6420
CACTTTGTAG	GGGTAGACAA	AGGGATTGTG	CGAACGAATG	TACTGACTGG	TAAAAACTA	6480
TCTGATAAGA	TTAGGAAAGA	AGTGGGCTCT	GGAGATAGCA	AACTAGGAAA	AGGCGGCTAT	6540
TTCTCTACTG	GGGATGTTCT	ATTAGGAAAA	GATGTTGTTT	CTTATACCGT	ACAAGTATTT	6600
TCAGAGAATA	ATGAAAGAGT	AGGAGTAAAC	ACTCAAAGTC	ACCGTGTTC	GTATAATCTC	6660
CCAATTCTAG	CTGACTTTTC	AGTCATCCAA	GATACTGTGG	AACCATCACG	AACCGTTGTT	6720
GAAAAATCA	TTCCAAAAC	AAATATTC	GAAGAAGAGA	AAGGGAAAA	AACCGAAGAA	6780
ATCAAGAAAA	AGAAAAAAC	CTCAGAATTG	GCAGAACTAA	TCTCAGAAAA	TGTGAAAGTT	6840
CGCTATGTTG	ATGAACAAGG	GCGTTTGCTA	TCATTGAAAA	ATGATACTGG	AATTGGAGAA	6900
AAAGAAAGTG	ACGGAACCTA	CATTACCAAT	AAAAACAAC	TGATTGGTAC	CAGCTATAAT	6960
GTCACAGATA	AAAAACTCAG	TAGCATGACT	ACTACTGACG	GAAAATATTA	TACTTTTAAA	7020
GAAGCAGATA	CAAATTCTGC	AAGTTTAACT	GGGAATATTG	TAAGCGAAGG	TAGAACAGTG	7080
ACCTTAGTTT	ATAGAGAAAG	CGAAGCGCCA	ACCACTGCTA	CAGTAACAGC	CAATTACTAT	7140
AAAGAAGGTA	GGCAAGAGAA	GTTGGTAGAG	TCTGTTATAA	AAGCTGATTT	AGCGATAGGT	7200
TCTGAGTATA	CCACAGAATC	AAAAACTATT	GAAGGGAAAA	CAACAACTGA	GGACAAAGAA	7260
GACCGAGTTA	TCACAAGGAA	AACAACATAC	ACCTTGGTAG	CAACTCCTGA	AAATGCGTAC	7320
CAGAAGACGG	TGCAACAGTT	GACTATTACT	ACCGTGAGAA	TGTTGAGGAA	ACAGTGGTTC	7380
CCAAACAGC	AACCTCTACT	GAGACGAAGA	CTATAACGCG	TATCATTCAT	TACGTTGATA	7440
AAGTTACGAA	CCAAAAATGTA	AAAGAAGATG	TGTTTCAACC	TGTAACCTTA	AGCCGTACAA	7500
AAACTGAGAA	CAAGGTCACG	GGAGTTGTAA	CCTACGGTGA	ATGGACAACA	GGAAACTGGG	7560
ACGAGGTTAT	ATCTGGTAAG	ATTGACAAGT	ACAAAGATCC	AGATATFCCA	ACAGTTGAAT	7620
CACAAGAAGT	TACGTCAGAC	TCTAGTGATA	AAGAAATAAC	GGTAAGGTAT	GACCGTTTAT	7680
CAACACCAGA	AAAACCAATC	CCACAACCAA	ATCCAGAGCA	TCCAAGTGTT	CCGACACCAA	7740
ACCCAGAACT	ACCAATCAA	GAGACTCCAA	CACCAGATAA	ACCAACTCCA	GAACCAGGTA	7800
CTCCAAAAAC	TGAAACTCCA	GTGAATCCAG	ACCCAGAAGT	TCCGACTTAT	GAGACAGGTA	7860

858

AGAGAGAGGA ATTGCCAAAC ACAGGTACAG AAGCTAATGC TACCTTGGCT AGTGCTGGTA 7920
 TCATGACCTT GTTAGCTGGT CTAGGATTAG GATTTTCAA GAAAAAGAA GATGAAAAAT 7980
 AATAGATTTT AGAATCTAGG AACCAGGAAA AGCTCACAGA TGTGGGCTTT TTTCCTGGTT 8040
 TTGAGAACGA GGTCTTTCGT AAAGAATAAA AACCTTACA AGTCTGTTGA ACTGGGAAAC 8100
 TATGAATCCT ATTTTTTTAA AAATATTTCC AGAAATCAGT TGCGG 8145

(2) INFORMATION FOR SEQ ID NO: 123:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 8697 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 123:

CGGTACCGGG AACGATACTT AGTCTAATTT TGCACCTTTT CCATGTATGG TAAAGGTTTT 60
 TCTTTTTTTA AAAAGGAAAA CGAGAAGAGG AGGTTCTTAT GAAAGCAAGC ATTGCCTTGC 120
 AAGTTTTACC CCTAGTACAG GGGATTGATC GGATAGCTGT TATGATCAG GTCATTGCTT 180
 ATCTGCawAC TCAAGAAGTG ACGATGGTAG TGACACCATT TGAAACGGTC TTGGAAGGGG 240
 AGTTTGATGA GCTTATGCGC ATTCTAAAAG AAGCGCTGGA AGTGGCAGGG CAGGAGGCAG 300
 ACAATGTCTT TGCCAATGTC AAAATAAATG TAGGAGAGAT TTTAAGTATT GATGAGAAAC 360
 TTGAGAAGTA TACTGAGACG ACACATTAGT CTATTGGGCT TTCTCGGAGT ATTGTCAATC 420
 TGGCAGTTAG CAGGTTTTCT TAAACTTCTC CCAAGTTTA TCCTGCCGAC ACCTCTTGAA 480
 ATTCTCCAGC CCTTTGTTCG TGACAGAGAA TTTCTCTGGC ACCATAGCTG GGCACCTTG 540
 AGAGTGGCTT TACTGGGGCT GATTTTGGGA GTTTTGATG CCTGTCTTAT GGCTGTGCTC 600
 ATGGATAGTT TGACTTGGCT CAATGACCTG ATTTACCCTA TGATGGTGGT CATTCAGACC 660
 ATPCCGACCA TTGCCATAGC TCCTATCCTG GTCTTGTGGC TAGGTTATGG GATTTTGCCC 720
 AAGATTGTCT TGATTATCTT AACGACAACC TTTCCCATCA TCGTTAGTAT TTTGGACGGT 780
 TTTAGGCATT GCGACAAGGA TATGCTGACC TTGTTTAGTC TGATGCGGGC CAAGCCTTGG 840
 CAAATCCTGT GGCATTTTAA AATCCCAGTT AGCCTGCCTT ACTTTTATGC AGGCTGAGG 900
 GTCAGTGTCT CCTACGCCTT TATCACAACT GTGGTATCTG AGTGGTTGGG AGGTTTTGAA 960
 GGTCTTGGTG TTTATATGAT TCAGTCTAAA AAAGTGTTC AGTATGATAC CATGTTTGCC 1020
 ATTATTATTC TGGTGTGAT TATCAGTCTT TTGGGTATGA AGCTGGTCGA TATCAGTGAA 1080
 AAATATGTGA TTAAATGGAA ACGTTCGTAG AATTAGAATG TTTCTGAAAA AGAAAAGAGG 1140

AAATCAAAAT	GAAGAAAACA	TGGAAAGTGT	TTTTAACGCT	TGTAACAGCT	CTTGTAGCTG	1200
TTGTGCTTGT	GGCCTGTGGT	CAAGGAAC TG	CTTCTAAAGA	CAACAAAGAG	GCAGAACTTA	1260
AGAAGGTTGA	CTTTATCCTA	GACTGGACAC	CAAATACCAA	CCACACAGGG	CTTTATGTTG	1320
CCAAGGAAAA	AGGTATTTC	AAAGAAGCTG	GAGTGGATGT	TGATTGAAA	TTGCCACCAG	1380
AAGAAAGTTC	TTCTGACTTG	GTTATCAACG	GAAAGGCACC	ATTTGCAGTG	TATTTCCAAG	1440
ACTACATGGC	TAAGAAATTG	GAAAAAGGAG	CAGGAATCAC	TGCCGTTGCA	GCTATTGTTG	1500
AACACAATAC	ATCAGGAATC	ATCTCTCGTA	AATCTGATAA	TGTAAGCAGT	CCAAAAGACT	1560
TGGTTGGTAA	GAAATATGGG	ACATGGAATG	ACCCAAC TGA	ACTTGCTATG	TTGAAAACCT	1620
TGGTAGAATC	TCAAGGTGGA	GACTTTGAGA	AGGTTGAAAA	AGTACCAAAT	AACGACTCAA	1680
ACTCAATCAC	ACCGATTGCC	AATGGCGTCT	TTGATACTGC	TTGGATTTAC	TACGGTTGGG	1740
ATGGTATCCT	TGCTAAATCT	CAAGGTGTAG	ATGCTAACTT	CATGTA CTTG	AAAGACTATG	1800
TCAAGGAGTT	TGACTACTAT	TCACCAGTTA	TCATCGCAAA	CAACGACTAT	CTGAAAGATA	1860
ACAAAGAAGA	AGCTCGCAAA	GTCA TCCAAG	CCATCAAAAA	AGGCTACCAA	TATGCCATGG	1920
AACATCCAGA	AGAAGCTGCA	GATATTCTCA	TCAAGAATGC	ACCTGAACTC	AAGGAAAAAC	1980
GTGACTTTGT	CATCGAATCT	CAAAAATACT	TGTCAAAAGA	ATACGCAAGC	GACAAGGAAA	2040
AATGGGGTCA	ATTTGACGCA	GCTCGCTGGA	ATGCTTTCTA	CAAATGGGAT	AAAGAAAATG	2100
GTATCCTTAA	AGAAGACTTG	ACAGACAAAAG	GCTTCACCAA	CGAATTTGTG	AAATAATGAC	2160
AGAAATTAGA	CTAGAGCAGC	TCAGTTATGC	CTATGGTCAG	GAGAGGATTT	TAGAGGATAT	2220
CAACCTACAG	GTGACTTCAG	GCGAAGTGGT	TTCCATCCTA	GGCCAAGTG	GTGTTGGAAA	2280
GACCACCCTC	TTTAATCTAA	TCGCTGGGAT	TTTAGAAGTT	CAGTCAGGGA	GAATGTCTCT	2340
TGATGGTGAA	GAAAAATCCCA	AGGGGCGCGT	GAGTTATATG	TTGCAAAAGG	ATCTGCTCTT	2400
GGAGCACAAG	ACGGTGCTTG	GAAATATCAT	TCTGCCCTC	TTGATTCAA	AGGTGGATAA	2460
GGCAGAAGCT	ATTTCCCGAG	CGGATAAAAT	TCTTGCGACC	TTCCAGCTGA	CAGCTGTAAG	2520
AGACAAGTAT	CCTCATGAAC	TTAGCGGTGG	GATGCGCCAG	CGTGTAGCCT	TACTCCGGAC	2580
CTACCTTTTT	GGGCACAAGC	TCTTTCTCTT	AGATGAGGCC	TTTAGCGCCT	TGGATGAGAT	2640
GACAAAGATG	GA ACTCCACG	CTTGGTATCT	TGAGATTCAC	AAGCAGTTGC	AGCTAACAAAC	2700
CCTGATCATC	ACGCATAGTA	TTGAGGAGGC	CCTCAATCTC	AGCGACCGTA	TCTATATCTT	2760
GAAAAATCGC	CCTGGGCAGA	TTGTTTCAGA	AATTAAACTA	GATTGGTCTG	AAGATGAGGA	2820
CAAGGAAGTC	CAAAAGATTG	CCTACAAACG	TCAAATTTTG	GCGGAATTAG	GCTTAGATAA	2880

860

GTAGAAAAAT	AGGGAGTTGG	TGAAGATTAT	CCTTTACCAG	CGCCCTTTTT	CTTTAAAAA	2940
TGAGAAAAAT	TCGGTATAAT	AGTCAAACAA	GGTCAAGGTT	TAAAGAGAGA	GGTGGGTTTG	3000
TTATGAGATT	TAAAAATACA	TCGGATCATA	TTGAGGCCTA	CATCAAGGCG	ATTTTAGATC	3060
AATCTGGTAT	CGTGGAGTTG	CAACGGAGTC	AGTTGGCAGA	TACCTTTCAG	GTTGTTCCTA	3120
GTCAGATTAA	CTACGTGATC	AAGACACGCT	TTACGGAAAG	TAGAGGCTAC	TTGGTTGAAA	3180
GTAAGCGTGG	TGGCGGAGGC	TACATTTCGTA	TAGGACGGAT	TGAGTTTCT	AGTCATCATG	3240
AAATGCTCCG	GGAGCTGCTT	TACTCGATTG	GTGAGCGAGT	CAGTCAAGAA	ATTTATGAGG	3300
ATATTCCTCA	GCTTTTGGTT	GAGCAGGAAT	TGATGACCAA	GCAGGAGATG	AATTTGCTAG	3360
AATCAGTAGC	TTTGGATCGC	GTTTTAGGAG	AAGAAGCTCC	AGTTGTTCGA	GCAAACATGC	3420
TACGTCAGAT	CATACAAGAG	GTAGATAGAA	AAGGGAAGTA	AGATGAACTA	TTCAAAAGCA	3480
TTGAATGAAT	GTATCGAAAAG	TGCCTACATG	GTTGCTGGAC	ATTTTGGAGC	TCGTTATCTA	3540
GAGTCGTGGC	ACTTGTGTGAT	TGCCATGTCT	AATCACAGTT	ATAGTGTAGC	AGGGGCAACT	3600
TTAAATGATT	ATCCGTATGA	GATGGACCGT	TTAGAAGAGG	TGGCTTTGGA	ACTGACTGAA	3660
ACGGACTATA	GCCAGGATGA	AACCTTACG	GAATTGCCGT	TCTCCCGTCG	TTTGCAGGTT	3720
CTTTTGTGATG	AAGCAGAGTA	TGTAGCGTCA	GTGGTCCATG	CTAAGGTACT	AGGGACAGAG	3780
CACGTCCTCT	ATGCGATTTT	GCATGATAGC	AATGCCTTGG	CGACTCGTAT	CTTGGAGAGG	3840
GCTGGTTTTT	CTTATGAAGA	CAAGAAAGAT	CAGGTCAAGA	TTGCTGCTCT	TCGTCGAAAT	3900
TTAGAAGAAC	GGGCAGGCTG	GACTCGTGAA	GATCTCAAGG	CTTTACGCCA	ACGCCATCGT	3960
ACAGTAGCTG	ACAAGCAAAA	TTCTATGGCC	AATATGATGG	GCATGCCGCA	GACTCCTAGT	4020
GGTGGTCTCG	AGGATTATAC	GCATGATTTG	ACAGAGCAAG	CGCGTTCTGG	CAAGTTAGAA	4080
CCAGTCATCG	GTCGGGACAA	GGAAATCTCA	CGTATGATTC	AAATCTTGAG	CCGGAAGACT	4140
AAGAACAACC	CTGTCTTGGT	TGGGGATGCT	GGTGTCCGGA	AAACAGCTCT	GGCGCTTGGT	4200
CTTGCCCAGC	GTATTGCTAG	TGGTGACGTG	CCTGCGGAAA	TGGCTAAGAT	GCGCGTGTTA	4260
GAACTTGATT	TGATGAATGT	CGTTGCAGGG	ACACGCTTCC	GTGGTGACTT	TGAAGAACGC	4320
ATGAATAATA	TCATCAAGGA	TATTGAAGAA	GATGGCCAAG	TCATCCTCTT	TATCGATGAA	4380
CTCCACACCA	TCATGGGTTC	TGGTAGCGGG	ATTGATTCGA	CTCTGGATGC	GGCCAATATC	4440
TTGAAACCAG	CCTTGGCGCG	TGGAACTTTG	AGAACGGTTG	GTGCCACTAC	TCAGGAAGAA	4500
TATCAAAAAC	ATATCGAAAA	AGATGCGGCA	CTTTCCTCGTC	GTTTCGCTAA	AGTGACGATT	4560
GAAGAACCAA	GTGTGGCAGA	TAGTATGACT	ATTTTACAAG	GTTTGAAGGC	GACTTATGAG	4620
AAACATCACC	GTGTACAAAT	CACAGATGAA	GCGGTTGAAA	CAGCGGTTAA	GATGGCTCAT	4680

CGTTATTTAA	CCAGTCGTCA	CTTGCCAGAC	TCTGCTATCG	ATCTCTTGA	TGAGGCGGCA	4740
GCAACAGTGC	AAAATAAGGC	AAAGCATGTA	AAAGCAGACG	ATTCAGATTT	GAGTCCAGCT	4800
GACAAGGCC	TGATGGATGG	CAAGTGAAA	CAGGCAGCCC	AGCTAATCGC	AAAAGAAGAG	4860
GAAGTACCTG	TCTACAAAGA	CTTGGTGACA	GAGTCTGATA	TTTTGACCAC	CTTGAGTCGC	4920
TTGTCAGGAA	TCCCAGTTCA	AAAAC TGACT	CAAACGGATG	CTAAGAAGTA	TTTAAATCTT	4980
GAAGCAGAAC	TCCATAAACG	GGTTATCGGT	CAAGATCAAG	CTGTTTCAAG	CATTAGCCGT	5040
GCCATTCGCC	GCAACCAGTC	AGGGATTCGC	AGTCATAAGC	GTCCGATTGG	TTCTTTATG	5100
TTCTTAGGGC	CTACAGGTGT	CGGGAAAAC T	GAATTAGCCA	AGGCTCTGGC	AGAAGTCTT	5160
TTTGACGACG	AATCAGCCCT	TATCCGCTTT	GATATGAGTG	AGTATATGGA	GAAATTTGCA	5220
GCTAGTCGTC	TCAACGGAGC	TCCTCCAGGC	TATGTAGGAT	ATGAAGAAGG	TGGGGAGTTG	5280
ACAGAGAAGG	TTCGCAATAA	ACCCTATTCC	GTTCTCCTCT	TTGATGAGGT	AGAGAAGGCC	5340
CACCCAGATA	TCTTTAATGT	TCTCTFGCAG	GTTCTGGATG	ACGGTGTCTT	GACAGATAGC	5400
AAGGGACGCA	AGGTCGATTT	TTCAAATACC	ATTATCATT A	TGACATCGAA	TCTAGGTGCG	5460
ACTGCCCTTC	GTGATGATAA	GACTGTTGGT	TTTGGGGCTA	AGGATATTCG	TTTTGACCAG	5520
GAAAATATGG	AAAACGCAT	GTTTGAAGAA	CTGAAAAAAG	CTTATAGACC	GGAATTCATC	5580
AACCGTATTG	ATGAGAAGGT	GGTCTCCAT	AGCCTATCTA	GTGATCATAT	GCAGGAAGTG	5640
GTGAAGATTA	TGGTCAAGCC	TTTAGTGGCA	AGTTTGACTG	AAAAAGGCAT	TGACTTGAAA	5700
TTACAAGCTT	CAGTCTGAA	ATTGTTAGCA	AATCAAGGAT	ATGACCCAGA	GATGGGAGCT	5760
CGCCCACTTC	GCAGAACCCT	GCAAACAGAA	GTGGAGGACA	AGTTGGCAGA	ACTTCTTCTC	5820
AAGGGAGATT	TAGTGGCAGG	CAGCACACTT	AAGATGGTG	TCAAAGCAGG	CCAGTTAAAA	5880
TTTGATATTG	CATAAAAGAA	TAAAAGTATC	AGCATCTGAC	CATAAGTCAC	AGTGGAGTGA	5940
AATCAATGA	AAATCAAAGA	GCAAAC TAGG	CAGCTAGCCG	CAGGTTGCTC	AAAACACTGG	6000
TTTGAGGTTG	CAGATAGAGC	TGACGTGGTT	TGAAGAGATT	TTCGAAGAGT	ATGAAACTAA	6060
AACCTATAGC	TTCTAAACGA	TCCGTGGTTT	TCATCAT TCA	ACACAAAATT	CATATGTTTA	6120
TTACCCCTCCG	TCGTATTTGT	CTTAGAGCGT	GTGTAGTAGA	AAAAGAGCAG	TCTTATCTGA	6180
AATTTTTATT	CTTTCAAAG	AGACCTGTTT	CTTTTTTGCA	TGTCAAATCC	GTTCTAGCTG	6240
GTATTTGAAA	AATCAAAC TA	ATATTCAATG	AAAATCAAAG	AACAAACTAG	GAAGCTAGCC	6300
GCAGGTTGCT	CAAAACACTG	TTTTGAGGTT	GTAGATAGAG	CTGACGTGGT	TTGAAGAGAT	6360
TTTCGAAGAG	TATAAGCTGC	AAGATGAATG	ATTTTCTTGT	ATTGACGTTG	TTGTTGACAA	6420

AAAGTAGCGG ATAAATGAAA TCCATTCCAT TATCATAGAT GATAGGCTGG TAGGAAATTT 6480
 TCAAATAGCA TACAGGAAAT AGATGTATGG AGTTCTGGTA GTAGAAAGGG AGAGAGATGA 6540
 ACATTTTAGT TGCAGATGAC GAGGAAATGA TTAGAGAAGG AATTGCAGCA TTTCTGACAG 6600
 AAGAGGGTTA TCATGTCATT ATGGCTAAGG ATGGACAAGA GGTCTTGAA AAATTTCAAG 6660
 ATCTCCCTAT CCATCTCATG GTACTGGATT TAATGATGCC TAGGAAGAGT GGTTTTGAAG 6720
 TGTTAAAAGA AATCAATCAA AAGCACGATA TTCCTGTCAT CGTCTTGAGT GCTCTGGGAG 6780
 ATGAAACTAC TCAGTCACAG GTATTTGATC TCTATGCTGA TGATCATGTG ACAAACCTT 6840
 TTTCTTTGGT ACTGCTTGTC AAGCGTATTA AGGCGCTTAT CAGACGTTAC TACGTCATAG 6900
 AGGATCTTTG GCGATATCAG GATGTAACAG TGGATTTTAC CTCTTACAAA GCACATTATA 6960
 AAAATGAAGA AATTGATCTC AAACCAAAGG AATTACTGGT ACTAAAGTGT TTGATTACAGC 7020
 ATAAAAATCA AGTTTTAAGT AGAGAGCAGA TATTGGAAGA AATTTCAAAA GATGTAGCTG 7080
 ATTTACCTTG TGATAGGGTC GTTGTATGCT ATATTCCGTAC TCTTCGCAAA AAATTAGCTT 7140
 TAGATTGTAT CGTGACTGTG AAAAATGTTG GGTATAAGAT TAGCTTATGA TAAAAAATCC 7200
 TAAATTATTA ACCAAGTCTT TTTTAAGAAG TTTTGCAATT CTAGGTGGTG TTGGTCTAGT 7260
 CATTCAATATA GCTATTTATT TGACCTTTCC TTTTTATTAT ATTCAACTGG AGGGGGAAAA 7320
 GTTTAATGAG AGCGCAAGAG TGTTTACGGA GTATTTAAAG ACTAAGACAT CTGATGAAAT 7380
 TCCAAGCTTA CTCCAGTCTT ATTCAAAGTC CTTGACCATA TCTGCTCACC TTAAGAGAGA 7440
 TATTGTAGAT AAGCGGCTCC CTCTTGTGCA TGACTTGGAT ATTAAAGATG GAAAGCTATC 7500
 AAATTATATC GTGATGTTAG ATATGTCTGT TAGTACAGCA GATGGTAAAC AGGTAACCGT 7560
 GCAATTTGTT CACGGGGTGG ATGTCTACAA AGAAGCAAAG AATATTTTGC TTTTGTATCT 7620
 CCCATATACA TTTTTGGTTA CAATTGCTTT TTCCTTTGTT TTTTCTTATT TTTATACTAA 7680
 ACGCTTGCTC AATCCTCTTT TTTACATTTT AGAAGTGAAT AGTAAAATGC AAGATTTGGA 7740
 TGACAATATT CGTTTTGATG AAAGTAGGAA AGATGAAGTT GGTGAAGTTG GAAAACAGAT 7800
 TAATGGTATG TATGAGCACT TGTTGAAGGT TATTTATGAG TTGGAAAGTC GTAATGAGCA 7860
 AATTGTAAAA TTGCAAAATC AAAAGGTTTC CTTTGTCCGC GGAGCATCAC ATGAGTTGAA 7920
 AACCCCTTTA GCCAGTCTTA GAATTATCCT AGAGAATATG CAGCATAATA TTGGAGATTA 7980
 CAAAGATCAT CAAAATATA TTGCAAAGAG TATAAATAAG ATTGACCAGA TGAGCCACTT 8040
 ATTAGAAGAA GFACTGGAGT CTTCTAAATT CCAAGAGTGG ACAGAGTGTC GTGAGACCTT 8100
 GACTGTTAAG CCAGTTTTAG TAGATATTTT ATCACGTTAT CAAGAATTAG CTCATTCAAT 8160
 AAGTGTACA ATTGAAAATC AATTGACAGA TGCTACCAGG GTCGTCATGA GTCTTAGGGC 8220

863

ATTGGATAAG GTTTTGACAA ACCTGATTAG TAATGCAATT AAATATTCAG ATAAAAATGG 8280
 GCGTGTAATC ATATCCGAGC AAGATGGCTA TCTCTCTATC AAAAATACAT GTGCGCCTCT 8340
 AAGTGACCAA GAACTAGAAC ATTTATTTGA TATATTCTAT CATTCTCAA TCGTGACAGA 8400
 TAAGGATGAA AGTTCCGGTT TGGGTCTTTA CATTGTGAAT AATATTTTAG AAAGCTATCA 8460
 AATGGATTAT AGTTTTCTCC CTTATGAACA CGGTATGGAA TTTAAGATTA GCTTGTAGAC 8520
 AGATTAGTTT TTTATTAAAG TTCATATAGG GTTAACATAA GTGTGTTATT CTTTGTGTAG 8580
 ATAAAAGAAA GGATACTAAT ATGGTATTAG CGATTATTTT AGTAACATTC TTTATTTCGAT 8640
 TGATTTTTTTT AAAGCGTTCG ATAGAGAATG AGAAACGAAT CCTTAGCAAT GCGCGGG 8697

(2) INFORMATION FOR SEQ ID NO: 124:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 4317 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 124:

AACCATACAT ACGGCAAGGC AAAGCTGACG CGGTTTGAAG AGATTTTCGA AGAGTATTAG 60
 TTGCCTTTAA AGGCATCCAC CATCGTTTGA AATCTTTCAT TTGAGAGAGT AATCCCTTTG 120
 CCCATTTTAG TATGGTCTGG ACTCCAAGCA CGAATATCAA ACTTTGCAGG GGCACCATTA 180
 AAGCTCACAC GGTAAATTC CTTGGTCCAA CCTTTTTTCGT TTTCAGAAAAG AGTCAACAAG 240
 TGCTCTTCGA TTTCAAATGT AAATTCGACC ATTTTCTTCT CCTTTTTTAG TTTCATTAGT 300
 TTATTCGTAA AATCTGTAG ATTTTAGGAA AATTTTATAT AATATTGATA TAAAAGAAGG 360
 GAGGCCAATA TGAGACATAA ATTCCAGCAA GTTCTAAATA AAATACATGA TTTTTTAAAT 420
 GGATATGACC AACCTGACCA GACTGAAACC AACTCCCTTA CAGCCACTAT TGAAGAGGCT 480
 ATCCAGAAAC AAACCGCTGT TCACCTTATC TTGTCTGAGA CAAGCTTTAC AGGTGACATC 540
 ATCAAATATG ATCAGCAAGG CCAGCAAATT ATCGTGAAAA ATTTTTCCAA AAATGTGAGC 600
 CGGATTATCC GTATAAGCGA TATTCAACGC CTGCGATTTG TCCCCTCAAC TGTCCAAACA 660
 GCCCAAAAAA ATAGATTTAA GAAAGAGTGA GATGTAGTTG CTTTATCCCA CTCTTTTTTC 720
 TTAGCGAATT TGTTCAAAAT GTAAATGAAC TGCGATATGA TCTCCATAAC CACTTCTTTC 780
 CAAGTCACGT TGTAACGAT AGGAAATGTA GTGTTCTGCA ATGGTAATGT AACCTGCGCC 840
 CAATAAACGA TGTTCAACCA TAGATTGAAT CATACTGATA GTCGCACGTT CCACCTTGCC 900

TTCTTGTA AAA TCCAAA ACTA CCTTCTTAGT GACTTGAGCA AGATTTTGAC GCAAATCATC 960
 TGTCAAAACA TAAACAGTTT GGGCTGCCTT CAAGATGGCT TGGTAAATCT TATCTGGATT 1020
 AAATTCAGCA ATTCGCCAT TACGTTTGAT TACTTGCA TA GGTTCCTCCT TTATTCCTTG 1080
 TTTTCTTTGA TTTCTGCCAG CATT TTTTCT TCTTCTACTG TCAGTTGATA ATGTTCAAGT 1140
 AAATCCGGTC TGCCTCGTA GGTTCCTTTT AACTCTCGT ACAATCGCCA CTGACGAATC 1200
 TTTTCATGGT GGCACACTCAT CAATACATCT GGCACGACCA TGCCTCGATA ATCATAGGGA 1260
 CGTGTGTACT GAGGATATTC TAAAAGACCT GAAGAAAAAC TATCATCTTG GTGGCTAGAC 1320
 TCCTTGCCAA TCACTTCTGG AATCAGGCGA ACTGTAGCAT CAATCATGGT CATAGCTGCC 1380
 AATTCCTCCAC CAGTGAGGAC ATAGTCACCT AGGAAATCT CATCTGTAC CAAGTCTTA 1440
 ATGCGCTCAT CATAACCCTC ATAGTGCCCA CAGATAAAGA TTAGCTCTTC CTCTTGAGCC 1500
 AAATCTTCAG CATAAGCCTG ATCAAACCTGC TTTCCAGCAG GATCAAGGAG AATAACGCGC 1560
 GGATTTTCT TTTCAATAGC ATCAAAGGAA TCGAAAATAG GTTGTGCTCT GAGCAACATG 1620
 CCCTGACCGC CTCCTAGGG CTCATCATCT ACATGACGGG CCTTTTCAGC ATTTTCTCGA 1680
 AAATTATGAT ACTGGATATC CAAGAGCCCT TTTTCTCGAG CCTTCCAAC GATTGAGTGC 1740
 TCCAGTGGAG AAAACATCTC TGGAAAGAGG GTTAAAATAT CAATCTTCAT CGTCTAACCC 1800
 TTCTAAGATT TCCACATCGA CCCGTTTACT TGGAAATATCA ACATTGAGAA CCACTGGTGG 1860
 GATATAAGGT AAAAGCAAAT CACGTTTGCC TTTTCGTTG ACCACCCAGA CATCATTAGC 1920
 ACCTGGTTGC AGGATTTCTT TGATGGTTCC AACCAAGCTA TCACCCTCAT AGACTTCCAA 1980
 ACCGATAATC TCGTGATAGT AAAATTCACC ATCGTCTAGG TCATTCAAAT CTTCTCAGC 2040
 GACCTTGAGA CTGTATCCCT TGTACTTTTC GATAGTATG ATATGGTACA TATCTTTGAA 2100
 TTTAATAATG TCAAAGTTCT TCTGTTTACG GTGGCTAGCG ATGGTCACTG TTTGGACAAA 2160
 CTGATCTTTT TCATCAAACA AAACCAGCTC AGCTCCTTTT TTAACCGTT CTTCTGCAA 2220
 ATCCGTCACA GACAAGACTC GCATCTCCCC CTGTAATCCC TCGTATTAA CGATTTTCCC 2280
 AACATTAAAG TAGTTCATCT TGTCTCCTGT AATCTCCTTT TTTCCATCTT ATTCTAACAA 2340
 TTCTCGAATA ATAGCCGCAA TTTTTCGGA TTCTGACCAT TGAAATAAT GGTGATTTCC 2400
 TCCTAAAATG AGTTTAGTAT TGAAGTCCA ATATCTGAT TCTCTGACT CTTTTCTCT 2460
 ATAAGGCTGA CAAAAACAA ATACAGGAAT ATGAGCTTCT ATAGATACAT CCTCAAAATC 2520
 TTCCTCAGTA ATCTCTCCAG ATATCTGAAA TTCTGGATCT TGATTTTCCA ACTCTAAGCC 2580
 TTTTCTTGC ATTAATCCC AGATTTTTTT ATTCGTTTCA GACTAAATG TTGCTTGAGT 2640
 TAAGTTCTTA AAATAAGTT CAGGACCACA CTCGCAATC AGCCTCATCT GCTCTCCAT 2700

TTCTGGATAA GGATTTTCTG AAAAATCAGC AAACATGACT TTTTTAGTTG TCGGTTCAAT 2760
 TGCTACTAAA GTCTGACGCT TAATTGGTTT CTCGAGTAAT TTGCAAGCTA AAATTCCTACT 2820
 CCAACTATGT GCACAAAGTA TATATTCAGA AATTCCTAAT TCTTCAAGTA CTTCATAAAC 2880
 CGCATCTGCA AGATTATCTA GATTTTTTCC AGCTTGGTCA TGAATCGGAC TCCTACCTGT 2940
 GTTCGGAAAA TCAATTGTCA AATAACCAAT TGTAGGAGGA GGTTTTTCAA GTATAAGTGA 3000
 AAAATTTTCA TAACTTGGTA GCAAACCTGC TCCGTTTAAA CAACTAGCA CTTTCTTTTG 3060
 CTTTTGATAA GTAACAGAGA GGCTACCAAT TTCTGTAGAT ACTTCAAACC TCTCATAAA 3120
 GAAATCCACT GATTCTATAT AATGAATTAT TAAAAATCCT TATCCTTTAT TTTATCACGT 3180
 TCCAAGGATT TTCTCAAGTT GGAGGAAGGG GACAATATCT CTTACTTTCCC TTCAATAATC 3240
 CTTCCAAAT ATGTTTATGT TGTAATTAA TGGCTGCGGT TTTGTCTTTC TCAAAGACAG 3300
 TCTTGGTAAG GTCAATATGA TTAATAGCTA CGATTGCGAC GGTGTAGTAA ATGATATCAG 3360
 CCAGTTCTCT GGCAAGTTCC TCCTTCGAAT CCTATCCCTT CTTTTCGACC AGAGCGCCTA 3420
 TTCAAAACCT CGACTACTTC TCCGACTTCC TCCACTAACT TCATAAAGAG ACCTTCATCA 3480
 GTCCGAGACT GCTGTTAATG TTCGATTAAG TAGTCTTGA ATTCCTTAAA CGTTCAATCT 3540
 TTTATAGTAT ATTGAAACTA GAATAGTACA CCTTACTTTC TAAAACATTG TTAGAAATCG 3600
 ATTTGACTGT CCTGATCGAT TTGTCCTGTT CTGTTTTCAT TTTACTATAT CTTCTATTCC 3660
 ACACAAAAAA GCGAGACATC CGTCCCGCCC TTCTTATTTT TCGTCAATAA CGATCTTTAC 3720
 TTTTTTGTAT TCAGTTGGGA CAGAGTAGAC AATCGTCTT ATCGCAGAAA TAGTGCAGACC 3780
 CTTACGACCG ATTACACGAC CCACATCGCT TTGATCAAGA TTCAAATGAT ATTCCAAAA 3840
 TTCTGGTGTA TCCTCAATCT TGATAGTTAA GGCATCTGGT TGTGAAATTA AGGGTTTCAC 3900
 AATCGCAATA ATGAGATTTT CAATCGTATC CATCTGTCAA CCTACTTTAA ACTTATTTTG 3960
 AAAATTTAGA ATCGTGAAT TTTTTCAATA CGCCTTCTTT TGAAAGGATG TTACGTA CTG 4020
 TGCTGAAGG TTGAGCTCCA TTAGCCAACC ATGCAAGAAC GCGGTCTTCT TTCAAAGTTA 4080
 CTTGGTTTTT AGCAACAAGT GGGTTGTAAG TTCCAAC TGTGATGAAA CGTCCGTCAC 4140
 GTGGTGAACG TGAATCTGCT ACGTTGATAC GGTAGAAAGG TTTTTTCTTA GAACCCATAC 4200
 GAGTCAAACG GATTTTAACT GCCATTTTAA AAGTCTCATT TCTTTAATTT TTTATTTCCG 4260
 TGAAATAGCT GAGCTATTTA GCACATGTTT TATTATAGCA GATTTCTGGC ATGTGTC 4317

(2) INFORMATION FOR SEQ ID NO: 125:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 4881 base pairs

866

(B) TYPE: nucleic acid
 (C) STRANDEDNESS: double
 (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 125:

AATTTATTTG ACTGGAAATT GTAGAGGGTT CTCGAAATTT CTTGAATGGT TAAAATAAGG	60
ACAAGAGAAA ACATGGATAT CTATATCCTT GTGCCAAAAA AACCACTGCC CTCCCAGAC	120
CAACCTGAGG AAAGCAGTGA TTCTTATTTT AGGAGTTAGG AATGAATACA CGAAATCAAT	180
TTAGCTGATT ATTTTTTGT TTTCAAGAAT TCATCGTATT GTTTTTGCAT TTCGTTCAAT	240
ACTTTTTCGT AGGCACCTTC AGATTTC AAT TTTCCATCA ATTCTGGAAT CGCTTTATCT	300
GGGTCTACAG TACCAGTGT GATAGCTGTA TCAAATTGTT GCATTGTGTT AGCAATAGCT	360
GAGATTTT CAG ATTTCACAT GTCAGTATTG AAGATAAATC CAAGCGCTGG AGATTCCTTTA	420
GCTTCTGCCA ATTCCTTCTT AGAATTTTCG ATTTGTTGGT CTGTAACGTT TTCGTTGATG	480
TAAAGGATCC AGTTGTTACC AGTGTTCAT CCACCCATGT GAGTGTTC TTTGTAGCCA	540
TCAAGAACGC GAACACGGTT TTCTTTACCT TCAATTTTTT CCCAGTTCTT GCCTTCTGGA	600
CCGTAAACAA GACCGTTCAA GAGTCTGGG TTCGTATTCA AGAGTTCAA GATTTCATT	660
GATTTTCTT TGTCTTAGA GTTGTGTTGAG ATGACAAAGT TAGCAACTTG TGTGTTTGG	720
TTTTTCTTGA TGAAGTAGT AATTGGTTT GATTGGATAT CTTTGTGGC AACACGTGAA	780
AGCAAGCTGT TACCGTAGTC AGCTGGTCCT ACTGTTTCTT CACGAACGAA CCAAGTATCT	840
TGTTGAAGGT CAAAGGAAGT ATCGCTTGT GCGACGCTT TTGGAATGTA GCCAGCTTCA	900
TAGAATTTGT GAAGAGTCTT CAAGTGTCTT TTGAAACGAG GCACTTCGTA ACGGTTTACA	960
ACTTTAGTAG TATCGCCTTC AAGTTCGATA ACGAATGGAA GACCGTTTGC TACTGGGTAG	1020
TCAAATTAT CAGATGGGAT GAAAACCTTA CCAATAGCAA ATGGTACTAC GTCTGGAGCT	1080
TTTTCTTTGA TTTGTTTCAA GACTGGCTCA AGAGTTTCGT AAGAAGTAAC ACCTGAAATA	1140
TCGATACCAT ATTTAGCAAG GAGAGTTCCG TTGAAGGCAA AGTTTGTAGA TGATGCAACG	1200
TTGGCTGCAA CTGGAACAGC GTAAATCTTA CCATTTACAG TATTACCCTT GATGTAAGCT	1260
GGTCAAGTG CTTTGTAAG GTCTTTACCT TCTTTTTTGT ACAATTCTGT CAAGTCAGCG	1320
TAAGCACCTT TTTGAGCATT TACAATATAG TTATCTGCAA AGGCAATATC ATAGTTTCA	1380
CCAGATGATG TGATAACTGA CATTTTCTTA CCATAGTCAC CCCAGCCAAG GTATTGGATA	1440
TCCAATTTGG CACCAACTTT TTCTTCAATG ATTTTGTGG CATTTGCTAA CAATTCATCC	1500
AGTTGTCTG GTTTGTCACC GATTGGTAC ATTTTGATAA CAGGTTTGT ACCTGAATCA	1560

GCAGCTTTTT	TGCTGTTACC	TGTCAAATTT	CCACAAGCAG	CAAGACCTGC	AGCCAGAGCG	1620
ACTACACTAG	CAGATGCAAA	AGCATATTTT	TTCCAGTTTT	TCATGATAAA	AACTCCTTTT	1680
TTTATTTT	AACTTATAAA	CAATGTAATG	ATCTTATACT	CAATAAAAAT	CAAAGAGCAA	1740
ACTAGAAAAC	TAGCCGCAGG	CTGCTCAAAG	CACTGCTTTG	AGGTTGTAGA	TAAGACTGAC	1800
GAAGTCAGTT	ACATATATCT	ACGGCAAGGC	GACGTTGACG	CGGTTTGAAT	TTGATTTTCG	1860
AAGAGTATTA	ACTTCACACA	AGGGAAGTTG	GGAAGTGAAG	AATGTTATTT	CTCAATAAGC	1920
ACTATCTTTT	CACACCACCG	ATAGTCAAAC	CTTTTACAAA	GTAGCGTTGG	AAAAATGGAT	1980
ACAAAATCGC	GATTGGAAGG	GTTGCAACCA	CAACCATGGC	CATACGACCT	GTTTCTTTTCG	2040
GTAGAGCAAC	TCCCAGTTGA	CCAATCAAGC	CGACCCTTTT	GGCAATGTAG	TCCATATTTT	2100
GTTGGATTTG	CATGAGCAAA	TATTGCAATG	GATACAAGTT	GTCACCTTTG	ATGTAAGAA	2160
GGGCGTTGAA	CCAGTCATTC	CAGAAACCAA	GAGCTGTTAA	GAGCGTGATG	GTTGCGATAC	2220
CTGGTAGTGA	CAATGGCAAA	CAGATTTGGA	AGAAAATCCG	GGCCTCACTG	GCACCATCGA	2280
TACGAGCCGA	TTCTAGAATG	GCTTCTGGAA	TGGTCTTCTT	GAAGAAGGAA	CGCATCAAGA	2340
TGATGTTAAA	TGGTGAGAGA	AGCATTTGAA	CAATCAAGGC	CCAAACAGTG	TCACCAAGCT	2400
GAAGTACACG	GGTCACCATG	ATATAACCTG	GTACCAAACC	AGCGTTGAAC	AACATACTGA	2460
GAAGGACGAA	GATGGTAAAG	AATCTGCGAT	ACTTAAAGGT	TGTCCGTGAA	ATAGCGTAGG	2520
CATAGGTTGT	TGTGATAAAG	ACATTTGTCA	ATGTCCCAAC	TACGGTTACA	AAGACAGAGA	2580
TGAAGAGGGC	TTGTAGGATT	TTATCTTAA	ACTGTGCCAA	AAACTCAAAA	CCGTCTAAGC	2640
CAAATTTGGA	TGGGAAGAAG	CTATAGCCGT	ATTGGAGGAG	GCTTTTCTCG	TCTGTCACTG	2700
AAATAATGAT	AACGAATACA	AAAGGTAGGA	TACAAGAGAG	GGCAATCAAA	CCCGAAATGA	2760
TACTGAAGAA	GATATCTGCT	TTCTTACTGA	AGGAGTGAAT	GCCGACATTA	TCAATTTTTT	2820
CTTTTTTAAT	TTTCTTTTTT	GCCATATTCT	CCTCCTTTCT	AGAACAAAGC	TGAGTTTGGA	2880
TCGACTCGTC	TTGCAAGCAA	GTTTGATAGG	ATAACCAGAA	TCAAACCAAC	AACGGATTGG	2940
TAAAGACCGG	CTGCTGCAGC	CATACCGATA	TCTGCTGTCT	GAGTCAAACC	ATTAAAGACA	3000
TATACGTCCA	AAACGTTGGT	TACATTGTA	AGCTGACCAG	CATTGTGTGG	GATTTGATAG	3060
AAGAGACCGA	AGTCTGCGCG	GAAGATATTT	CCGACTGCAA	GGATGGTCAA	TACAGTTACA	3120
AGCGGAGTCA	ACTGAGGAAT	GGTTACGTTG	CGAATACGTT	GCCACTTGCT	AGCTCCGTCC	3180
ACTGTCGCTG	CTTCGTAGTA	GGTTGGATCA	ATTCACATGA	TCGTCGCATA	GTACATGACA	3240
CTGCTATATC	CAAAGCCTTT	CCAAATACCT	AGGAAAAGTA	GGAGATAGGG	CCAGATGCCC	3300

868

AGGTCAGCGT AGAAATTGAC TTCTTTGAGA CCAAGACTTT CCAATAGATG ATTGAACACC 3360
 CCTTTATCAA TATTTAGGAA GGCATCTGTA AAGAACTGA TGATAACCCA AGACAAGAAG 3420
 TAAGGGAACA ACATAGAAGT TTGAAAAATC TTCACCATTC TCTTAGAACG GAGCTCGCTG 3480
 AGGATAATGG CAATCCCTAC AGATACAAC T AACCTAGAA AGATAAAGCC AAGATTGTAG 3540
 AGGACAGTAT TTCGTGTGAT AATAAAGCG TCTCTTGAAC TAAATAAGAA TCTAAAATTA 3600
 TCGAGTCCGA CCCATTTACT ATTTATGATA CTATCTATGA AACCATTACT GGTCATGTGG 3660
 TAGTCTTTGA AGGCAACCAC GTTCCCAAAT ACTGGAATGT AAAAGAATAG AATCAACCAG 3720
 AGTGCCCTG GCAAAACCAT CAAGAGAAAG ATCCAGTTGT CTCTCAATGT TTTTGAAAAC 3780
 TTTTTCATAA TTTCTCCCT TTTTATTTT ATATCCATCT AAAAAATCTT TTTTAGACTT 3840
 TTGATAACGA TTACATTAT AGTATACTCC TATTTGCAGG TTAGGTAAA CTCCTAATTA 3900
 TAGAAAAAC TCCACAAAT ATGTAGCAGA TTTAAAAC T TATCACCAC ATCAAACAAA 3960
 TGTCTAAAT CAATGTTTA TTTTATCTCT ATTAGCCCAG TGATGGCGTC ACTCTGTTAT 4020
 AAGCATCCAA CAACGGGTA TACTGAAAA TCTCCAGACT AGGGAECTA GCGATAGTTC 4080
 CTAATCTGGA GATTTTAAAT ATGTTATTAG GCGTTGCTT TCAACTTAGC AATAACCTCT 4140
 TTAAGATTAT CAATCAACTC TGCTGCAGTA TGCTCAGAGC CTTTTTCATC TGCCAAGAAC 4200
 AAAACTGCTT TTTGAAGTTC TTTTGTAGAG TTTTCAAGGA CATCCTTATC TACTGTTTCA 4260
 AGGTTTGAGT CTTAAGAAG TTTACTTAAT TCCTTGGCTA ATTTCTTGAG TTTGATTTGC 4320
 AGACTCATCT TCTCTGTG TTTCTTTGCC CGCTGTTTGT CCTCCATCCT TAGTTGCTGA 4380
 CTGGCTTTCC TTAATGGACT CTAGGGAAGC AATGGCATCT TTGACTGTTT GCAAGATATC 4440
 ACGTAAACCT TGCTCTGTCA AACTATCATC TGCAAAAGCT TTATTAGCCT CTGCCAAAAC 4500
 CAGACGTGCT GAATCTGTGG TAGGATTCGA TACACCTGTC AATGATCTCA AAAGATTTTC 4560
 TAAGGTTTGA GTCTGCTTAC TAATACTAGA CTAAAATCAA AAAGTATTAT ATAACAGTGA 4620
 TATGAAATCA ACTAAAGAAG AAATCCAAC CATCAAAACA CTTTAAAAG ACTCTCGTAC 4680
 AGCTAAATAT CATAAACGCC TTCAAATCGT TCTATTTTGT CTGATGGGCA AATCTTATAA 4740
 AGAGATTATA GAACTTTTAT AGTAGTTGA AATAAGATGT GAACATCTCT ATCAGGAAAG 4800
 TCAAATTAAT TTATAGAAAT ATTTTAGCAG CCAAGGTGTA CTGTTATAGA TTCAATACAC 4860
 TATACTGGT GGTTTAGCTC G 4881

(2) INFORMATION FOR SEQ ID NO: 126:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 13121 base pairs
 - (B) TYPE: nucleic acid

(C) STRANDEDNESS: double
 (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 126:

AGGATCCCCG	GAAAAGGAGA	CTAAAAATGA	AGAAAAAATT	TCTAGCATT	TTGCTAATTT	60
TATTCCCAAT	TTTCTCATTA	GGTATTGCCA	AAGCAGAAAC	GATTAAGATT	GTTTCTGATA	120
CCGCCTATGC	ACCTTTGAG	TTTAAAGATT	CAGATCAAAC	TTATAAAGGA	ATTGATGTTG	180
ACATTATTAA	CAAAGTCGCT	GAGATTAAAG	GCTGGAACAT	TCAGATGTCC	TATCCTGGAT	240
TTGACGCAGC	AGTCAATGCG	GTTCAAGCTG	GGCAAGCCGA	CGCTATCATG	GCAGGGATGA	300
CAAAGACTAA	AGAACGTGAA	AAAGTCTTCA	CCATGTCTGA	TACTTACTAT	GATACAAAAG	360
TTGTCAATGC	TACTACAAAG	TCACACAAAA	TTAGCAAGTA	CGACCAATTA	ACTGGCAAAA	420
CCGTGGTGT	TAAAAACGGA	ACTGCCGCTC	AACGTTTCCT	TGAAACAATC	AAAGATAAAT	480
ACGGCTTTAC	TATTAACA	TTTGACACTG	GTGATTTAAT	GAACAACAGC	TTGAGTGCTG	540
GTGCCATCGA	TGCCATGATG	GATGACAAAC	CTGTTATCGA	ATATGCCATT	AACCAAGGTC	600
AAGACCTCCA	TATTGAAATG	GATGGTGAAG	CTGTAGGAAG	TTTTGCTTTC	GGTGTGAAAA	660
AAGGAAGTAA	ATACGAGCAC	CTGGTFACTG	AATTTAACCA	AGCCTTGCTC	GAAATGAAAA	720
AAGATGGTAG	TCTTGATAAA	ATTATCAAGA	AATGGACTGC	TTCATCATCT	TCAGCAGTGC	780
CAACTACAAC	TACTCTCGCA	GGATTAAAAG	CTATTCCTGT	TAAGGCTAAA	TATATCATTG	840
CCAGCGATT	TTCTTTTGCC	CCTTTTGT	TCCAAAATTC	AAGCAACCAA	TACTCTGGTA	900
TTGATATGGA	ATTGATTAAG	GCAATCGCTA	AAGACCAAGG	TTTTGAAATT	GAAATCACCA	960
ACCCTGGTTT	TGATGCTGCT	ATCAGTGCTG	TCCAAGCTGG	TCAAGCCGAT	GGTATCATCG	1020
CTGGTATGTC	TGTCACAGAT	GCTCGTAAGG	CAACTTTTGA	CTTCTCAGAA	TCATACTACA	1080
CTGCTAATAC	CATCTTGGT	GTCAAAGAAT	CAAGCAATAT	TGCTTCTTAT	GAAGATCTAA	1140
AAGGAAAGAC	AGTCGGTGT	AAAAACGGAA	CTGCTTCTCA	AACCTTCCTA	ACAGAAAATC	1200
AAAGCAAATA	CGGCTACAAA	ATCAAACCT	TTGCTGATGG	TTCTTCAATG	TATGACAGTT	1260
TAAACACTGG	TGCCATTGAT	GCCGTTATGG	ATGATGAACC	TGTTCTCAA	TATTCTATCA	1320
GCCAAGGTCA	AAAATTGAAA	ACTCCAATCT	CTGGAACTCC	AATCGGTGAA	ACAGCCTTTG	1380
CCGTTAAAA	AGGAGCAAAT	CCAGAACTGA	TTGAAATGTT	CAACAACGGA	CTTGCAAACC	1440
TTAAAGCAAA	CGGTGAATTC	CAAAAGATTC	TTGACAAATA	CCTAGCTAGC	GAATCTTCAA	1500
CTGCTTCAAC	AAGTACTGTT	GACGAAACAA	CGCTCTGGGG	CTTGCTTCAA	AACAACCTACA	1560

870

AACAACCTCCT TAGCGGTCTT GGTATCACTC TTGCTCTAGC TCTTATCTCA TTTGCTATTG	1620
CCATTGTCAT CGGAATTATC TTCGGTATGT TTAGCGTTAG CCCATACAAA TCTCTTCGCG	1680
TCATCTCTGA GATTTTCGTT GACGTTATTC GTGGTATPCC ATTGATGATT CTTGCAGCCT	1740
TCATCTTCTG GGAATTCCTA AACTTCATCG AGTCTATCAC AGGCCAACAA AGCCCAATTA	1800
ACGACTTTGT AGCTGGAACC ATTGCCCTCT CACTCAATGC GGCTGCTTAT ATCGCTGAAA	1860
TCGTTTCGTTG TGGTATTCAG GCCGTTCAG TTGGCCAAAT GGAAGCCAGC CGAAGCTTGG	1920
GTATCTCTTA TGAAAAACC ATGCGTAAGA TTATCTTGCC ACAAGCAACT AAATTGATGT	1980
TGCCAAACTT TGTC AACCAA TTCGTTATCG CTCTTAAAGA TACAACATC GTATCTGCTA	2040
TCGGTTTGGT TGAACCTCTC CAAACTGGTA AGATTATCAT TGCTCGTAAC TACCAAAGTT	2100
TCAAGATGTA TGCAATCCTT GCTATCTTCT ATCTTGTAAT TATCACACTT TTGACTAGAC	2160
TAGCGAAACG CTTAGAAAAG AGGATTTCGT AATGGCAAAA TTA AAAATG ATGTAAATGA	2220
TTTACACAAG CACTATGGAA AAAATGAAGT CCTAAAAGGA ATTACGACTA AGTTCTATGA	2280
AGGAGATGTT GTTTGTATCA TCGGTCCTTC AGGTTCTGGT AAGTCAACTT TCCTCCGTAG	2340
CCTCAATCTT TTAGAAGAAG TCACTAGCGG TCACATCACT GTGAACGGCT ATGATTTAAC	2400
TGAAAAACA ACCAATGTG ACCACGTCG TGAAAATATC GGCATGGTAT TCCAACACTT	2460
CAACCTCTTC CCTCATATGT CTGTATTGGA CAACATCACC TTTGCTCCTA TTGAGCACAA	2520
GTTGATGACT AAGGAAGAAG CTGAGGAATT GGAATGGAG TTGCTTGAAA AGGTTGGACT	2580
AGCAGATAAA GCTAATGCCA ATCCAGATAG CCTATCAGGT GGTCAAAAAC AACGTGTGGC	2640
CATCGCTCGT GGCCTAGCAA TGAATCCAGA CATCATGCTC TTCGATGAAC CAACTTCTGC	2700
CCTTGACCCT GAGATGGTTG GAGACGTACT TAACGTTATG AAGGAATTGG CTGAGCAAGG	2760
CATGACCATG ATTATCGTAA CCCATGAGAT GGGATTTGCT CGTCAGGTTG CCAACCGCGT	2820
TATCTTTACT GCAGATGGCG AGTTCCCTGA AGACGGAACA CCTGACCAA TCTTTGATAA	2880
CCCACAACAC CCTCGTCTGA AAGAGTTCTT AGATAAGGTC TTAAACGTCT AACTCAAAC	2940
TGTAAGGATT TCCTTGCACT TTTTCTACCT CGTATTGGAA TTTTGTGATT TTCGGAAAA	3000
TATGTTAGAA TTAAGTTTAT GAAATGAGGT TTCCTCATA C TAGCAAGAC TAGGAATAAA	3060
AATAGAAATT AGGTAGCTAG ATGTCATCTA AGGTTATTGT TACAATTTT GGTGCGAGTG	3120
GAGACCTGGC TAAACGCAAG CTCTACCCTT CCCTTTT TAG ACTATATCAA TCCGGCAATC	3180
TTTCCAAGCA CTTTGCCGTT ATTGGAACCT CCCGTAGACC TTGGAGTAAG GAATATTTTG	3240
AATCTGTAGT TGTCGAGTCC ATCCTTGATT TGGCAGATAG TACCGAGCAA GCCCAAGAAT	3300
TTGCTAGCCA CTTTACTAT CAAAGCCATG ATGTCAATGA TTCGGAACAT TATATTGCTT	3360

TGCGTCAATT ACAAGCTGAG CTTAATGAAA AATACCAAGC TGAACACAAT AAGCTCTTCT 3420
 TCTTGTCTAT GGCACCTCAG TTCTTTGGAA CCATTGCCAA ACACCTCAAA TCTGAAAACA 3480
 TTGTGCGATGG CAAAGGTTTT GAGCGCTTGA TCGTTGAAAA ACCATTTGGT ACAGATTACG 3540
 CAACTGCAAG CAAGTTGAAT GACGAACTCC TAGCAACATT TGACGAAGAA CAAATTTTCC 3600
 GTATCGACCA TTATCTTGGT AAGGAAATGA TCCAAAGCAT CTTTGCAGTT CGCTTTGCAA 3660
 ACTTGATTTT TGAAAACGTT TGGAAACAAG ATTTTATCGA CAATGTTCAA ATTACCTTTG 3720
 CGGAGCGCTT GGGTGTAGAA GAACGTGGTG GCTACTATGA CCAATCCGGT GCCCTCCGTG 3780
 ACATGGTCCA AAACCACACT CTACAAC TTCGCTCCT CGCCATGGAC AAACCAGCAA 3840
 GCTTCACAAA AGACGAGATT CGTGCTGAAA AGATTAAGGT CTTTAAAAAC CTCTATCATC 3900
 CAACTGATGA AGAACTCAAA GAACACTTTA TCCGTGGGCA ATACCGCTCT GGTAAAGATTG 3960
 ATGGCATGAA ATACATCTCT TATCGTAGCG AGCCAAATGT GAATCCAGAA TCAACAAC TG 4020
 AAACCTTTAC ATCTGGTGCC TTCTTTGTAG ACAGCGATCG ATTCCGTGGT GTTCCTTTCT 4080
 TTTTCCGTAC AGGTAAACGA CTGACTGAAA AAGGAACTCA TGTC AACATC GTCTTTAAAC 4140
 AAATGGATTC TATCTTTGGA GAACCACTTG CTCCAAATAT TTTGACCATC TATATTCAAC 4200
 CAACAGAAGG CTCTCTCTT AGCCTAAATG GGAAGCAAGT AGGAGAAGAA TTTAACTTGG 4260
 CTCCTAAC TC ACTTGATFAC CGTACAGATG CGACTGCAAC TGGTGCTTCT CCAGAACCAT 4320
 ACGAAAAAT GATTTATGAT GTCCTAAATA ACAACTCAAC TAACTTTAGC CACTGGGATG 4380
 AAGTTTGTGC GTCATGGAAG TTGATTGACC GTATTGAAAA GCTCTGGGCT GAAAATGGTG 4440
 CCCCAC TTCA T GACTATAAAA GCTGGAAGCA TGGGACCTCA AGCCAGCTTT GACCTACTTG 4500
 AAAAATTCGG TGCCAAATGG ACTTGGCAAC CAGATATCAC CTATCGTCAA GATGGTCGCT 4560
 TAGAATAAAA AAATTTCTG CAAGTTTATG CcTTGCAGGA TTTTGTCTTC TGATTAGATT 4620
 AAACCTTCCA AGAGACCTTT CATAAAGTTT TCTGAGTTAA ACTCTCCAAT ATCATCGATT 4680
 TTTTACCAA AACCAATCAA TTTTACAGGA ATATTGAGTT CTTCACGAAT GGCTAGAACC 4740
 ACACCTCCTC GAGCAGTTCC ATCAATCTTA GTCAAACAA TTCCCGTTAA AGGTGTGATT 4800
 TFCGAAAATT CTTTGGCCTG TACTAGGGCA TTTTGACCTG TTGATGCATC AAGTGCCAAG 4860
 AAGGTTTCAT GTGGTGCTTC TGGCACAACA CGTTTGATAA TACGACCAAT CTTTTC AAC 4920
 TCAGCCATAA GGTTATCCTT ATTTTGCAGA CGACCAGCAG TATCAATCAT GAGAATATCG 4980
 ATACCTTCAG TCACGGCAGC TTCCATACCA TCAAAGACCA CGCTGGCTGG ATCAGCTTTT 5040
 TCAGGTCCAG TTACTACTGG AACATCTACT CGTCGGCCCC ATTCAGCTAG CTGAGCTACT 5100

872

GCACCCGCAC GGAAGGTATC TGCTGCAACC AGCATGACCT TCTTACCAGC TTGTTTGTAG 5160
 CGGTGGGCTA GTTTTCGGAT AGAAGTTGTT TTCCCAACAC CATTACACACC AACAAAGAGC 5220
 ATAAGTGTCA AGTTATCTTG GAAGTGGATG CTTTCATCGT AGCTACCATC CTTTTCATAA 5280
 AGCTCAACCA ATTTCTCAAT GATGACACGA CGAAGTACAT CAGGTTTCTT GGCATTTTCA 5340
 AGCTTGGCTT CGTAACGTAG TTCTCCGTT AAGTTAGAAG CGACTTGGAC ACCAACATCA 5400
 CTCATAATCA GCAGTTCTTC CAGTTCCTCG AAAAATTCCTT CGTCAACAGA GCGGAAGTTA 5460
 GCAAAGAAGG CATTCAAGCG GGCACCGAAA CCTGTGCGAG TTTTCTTAAG ACTGCGGTCA 5520
 TATTTTTCCCT GAACAGTTTC TTCTGTTTGA GGAGCTTCTG GTTCAAGCAC TTCAGAATTA 5580
 TTTTCTTCTA CAGTTCCTTC GTGCTCAAGC TTCTCTTCCCT CTGGTAATTC TTCTGAGTTT 5640
 GGTAATTCCT CTATTTCTTC TTGAGAAACC CCTACAGCTG GCTCTGAATC CTGACTTTCT 5700
 TCAACTGTGT CTTGGATTTC CTCTTCTTGG AACACAGCTT GTTCAACAAT TTCAACCTCT 5760
 GCTTCTTCCCT GAGAAACTTC CTCAACTTCT GTGAAGGTAG GATCAACATC TTCAGACAAA 5820
 TCAAGATTTT CCAGAGCTTC TTTTACAAC TCTTCGATTT TAGGTTCTTC TTTTTTTCCG 5880
 AATAGACGGT CAAACAATCC CATATCTTAG TTCTCCTTTA GCACATATTC TTCGATAGCC 5940
 CAGGCGACAG CTTCTCCTATC GTTGGTCATC GCGCTCACTA CATTGCGGC TGCCTTTACT 6000
 TCAGGAACAG CGTTTTGCAT AGCAACACCA AGACCTGCCC ATTCAATCAT AGAGAGGTCA 6060
 TTGGCCTCGT CACCACAAGC CATCACTTGA CTTTGGTTCGA TTCCAAGATG GCTGATTAGT 6120
 TTTGCCAAAC CTGTTGCTTT ATGAACATTC TTTGGTGACC ATTCTAGCAA CATTTCACGT 6180
 GATTTAAAGA TTTTATATTG GTCAAACAAT TCTGGAGAAA TCTTCTGAAT GGCTGCATCC 6240
 AAGGGTTCTT GAGCAAAGGC AGTCACGCAT TTGTTGTAGG TCATTTGACT AGATAAGTCT 6300
 TCAAAGTCCA CTGGAACAAA GGTCAAAGCT GGATTGAATT TGGCATAAAG ACTTCTTGG 6360
 TCCGATTGGA TTTGATAAAC TGTTCCTTCT GAGATGGCAT CAAGAGGCAG TGATAATTTT 6420
 TCTGTTTCTT CATACAAACG TGCCACATCA TCATATGAAA AGACTGTTTT ATCAAGGATT 6480
 TCTCCTGTAT TTTTCTGAAC TAATCCACCA TTAAAAGTAA TGGTATACTC ATCTTCTGTA 6540
 CCGTCAGTCC CTAATCATG GAGAAAGAAA TCCATGGCTT TTAAGGGACG ACCAGTTGTC 6600
 AATACGACCT TGATACCACG ATCACGCGCA gCTTGCAAGG TTTCCCTGGT ACGATCCGTC 6660
 AGCCTTTTAT CAGTAGTCAG CAAGGTCCCG TCCAAGTCCA ATGCAATCAA TTTTATATCT 6720
 GCCATTATAA GCCCTCCATA TAAGCTATAA CCGACCGTTC CTTATGGTGA CCAATCACAG 6780
 TCTTTGCTAA TTCTAAAATT TCAGGTCGTG CATTTTCAGG AGCTACAGGA TGTCACCAA 6840
 CCTGCATCAT ATGTAAGTCA TTAAGATTGT CTCCAAAAGC CATGACCTGA TCCATTGTGA 6900

TACCAAGTTT TTTAACTAAT TCAACAATGG CCACTCCCTT ATCGACATAG TCCAGAACAA 6960
 TATCAATGGA TTCAAAGCCA GTTGTCATGG CCTTAACACC AGGAACGTTT TCGTTTACCC 7020
 AAGCCTCCCC ATCTTCCAGC GTTCTTCTG TGAAGTTGGT TGTAATTTG AAAATGTCAT 7080
 CTGTGATATC TTCCAAACTC GCTACTTTTT GGATATTTTC ATTATAGTGC TGACTIONCT 7140
 TCAAATAGGT CTCATCAACC GTATCTAGAA CATATGAACC CTCTTACCC GTCAAGAGCA 7200
 GTTATTGAT ATCTACATAA GGTGAAGTTT TCAGCTTTTC AAAAGTTGCC AGATAAAAAGT 7260
 CACGAGACAT AGTCGCTTCA TACAAGTCCT GACCTTGATA CTCTACCAA CTGCCATTTT 7320
 CCGCGATGAA AATAATGTCA TCACGAACAC CAGCAAATAA TTTTCTAGA GACAGAAATC 7380
 CCCGACCCGA AGCTACCGCA AAGTAAATCC CTTTTTCCTT GTAGGAAACC AAGAGAGACT 7440
 TGAGACGATC CATATCAAAG CGTCCATTCC CATCTAGGAA GGTCCCGTCC ATATCCGTTG 7500
 CTACTIONTTT AATTGTCATC CTCAATACT TTCTAAATCT TTTAACTTAA CTGAAACAA 7560
 CTTTGAAACA CCCGATTCTT GCATGGTCAC TCCATAGATG GAATCAGCCG CTGCCATGGT 7620
 TCCCTTACGG TGGGTTACGA CGATGAACTG GCTGTCTTTC TCAAAGCGGT TGAGGTAATC 7680
 CCCAAAACGT TTAACATTGG CTTCATCCAG CGCAGCTTCC ACCTCATCCA AGATAACAAA 7740
 TGGAATAGTC TTGACACGAA TAATGGAGAA GAGCAAGGCA AGAGCCGATA GGGCTTTTTT 7800
 ACCACCCTC ATGAGATTAA GAGACTGGAT TTTCTTGCCT GGTGGTTGGA CAGAAATTTT 7860
 AACCCAGCT GTCAGCAAGT CTCCTTCAGT CAAAATGAGG TCAGCCTGAC CTCCACAAA 7920
 CATCTGCTTG AAGGTCACTT TAAAGGACTC ACGAATGACC TCAAAGGTTG ATTTAAAGCG 7980
 TTCCTTGACC TCATCATFCA TCTCTGTAAT GGTCTCAAGG AGCAGGTTTT TCGCAGACAA 8040
 AATATCATCA CGTTGGCTAT TTAGGAAATC CAGACGGTTG TGAACCTTCTT CGTACTGTTC 8100
 AATAGCGTCT AAATTGACAG GACCCAGTGA GCGTATAGCC TTCTCTAAAT CCTTAACTTC 8160
 TTGCTCTGCC AGATTGAGAT TTTCCAACTC ATGCGCCTTT TCTAAAGCTT CTGTGTAGCT 8220
 GATCTGGTAC TGGTCTGTTA ATTGACTTTG TAGATGGCGC AAGCGCTCGC TAACCTTTTC 8280
 TTTCTTGGCT TCAGCACGAG TTTGCTTGCG AATCCACTCT TCATTCTGCT GCGGAGCCTG 8340
 ATCCAAATGA CTAGCAATAT CATCCAGTTG ACCCTCAATA TCATCCAACT CAAACTGCTT 8400
 GCGAATCAA CCTTGTGGA GATTTGTTTT TTGAGTTTTG GATTCTTCCG CCTGTTGACT 8460
 GAGCAATTCT GTATCAACCT TCTCAAGATT ATCAATCTTT TCTTGAAGAA GCGCTGGAT 8520
 TTCCTCTGT TCAAAATCAA GATTGTCAA TTCCTTGCTT AAGCGTTCAA TATCAGCAAC 8580
 TTCATAACGT TTTTGCCCTT GCAGTCTGT CTTAAGCAA CGAGCTTGCG CTAGCTCTTC 8640

CTGCAAGTTT TGATAGCGTT CTTGGATGGC ATTTTGTGTTA GACTTAATCT CTTCAATCTC 8700
 AGCTTCCAGA TTTTGCTTGT CACTGGAGAT TGCAGCAAGA CGCTCTTGGC AGTTTTCCTT 8760
 ATCCGCTTGC CAATCTCCCT CGGAAAGACG ATCTATTTCC TCTTCTTGGA GTTTCCAAAG 8820
 AGTTTCCAGT TCTTCAACTT GCTGACTAGT TTGCTGATAA GCGAGGAACA AGCCTTGCTC 8880
 CTGAATACGT GCCTGCTCTC CTTGAGATTT AATAGCTTCT AATGACTCGG TCAATCTGGC 8940
 CATCTCATCT TGCAAGGTCT TCAAAGTCGC CTCTTCTGAA CCCAAGCTTG CTCTTCTTC 9000
 AGCAATTTCT TTTTGTAATT GCTCCAGTTC TGGCTTGATA AAAATGCTGT TATTCTGGCG 9060
 ATTGGCACCA CCTGCATAAG AACCACCTGT GCGCAACTCT GTCCCATCCA ATGTCACCAT 9120
 ACGAACCTGA TAACGAACTT GCGGAGCTGC TGCACGCGCA TGTTCTACGG TATCAAAGAT 9180
 AGCCGTCGTA GCTAGCAAGT TCTTGAAAAT GGCTTCCAGT CTAGTATCAA AAGTCACCAA 9240
 CTCATCTGCC ATCCCAAGGA AACCTGGGCT TACAGCGATA GCATCTGGT TCTGACTAGA 9300
 AATCGTACGC GCCTTGATAG TGGTCAAAGG AAGAAAGGTT GCACGACCGG CTCTGTTCGG 9360
 TTTAAGGAAG TCAATAGCCT TGGTTGCCGA CTCTTCATCT TCTACGATGA TATGCTGGCT 9420
 ACTTGCCCTT AAGGCAATCT CTAGGGCAGT TTGATAATAA ACATCAAAGG TCAGATGCTC 9480
 ACTGACTGCA CCAATAATCC CACCTAGGCG ATCTTTTCTT TGGAGAACAC TCTTAACACC 9540
 TGCATAAAAG TTACTATGAT TTCTCAGGAT ATTTTCCAAA CTTTGAGCTC TGGCCTGCTT 9600
 GTTTTGTGAGA TTATCCAGAC GGTCAAAGAG TTGGCTTTGT TGAGCTTGAT AGGAAGTTT 9660
 CTGCTCCTCT TGCTCCTTGG CAATAGCTTG GTAGTCAGCC AATAATTCTT GAACCTGCTC 9720
 CTTGGCAGTT TCAAGCTCTT CCTTTTGCTG ACTAGCCTTC TCTTTAGCTA TAGCTAATTG 9780
 CTCTTTCAGC TTTTCTAGTT GATCTGCTTG TTTTGTGAGAA AGCTGACGAC TATTTTCCAA 9840
 CTCATTCTCA ATACGGGTCA ACTGGTTTGA GACATCCGCT TCTTCTTGTA AAAGAGCTAC 9900
 AAAGCGTTCA CGTAAGAGCT CAATCATCTG ATCAGGATCG TCTGAGAAAG CCAGCAATTC 9960
 AGCTTCTAAA CGATTGAGTT TTTGATTATT TTGGACTAGA TTTCCCTCTA ACAGAGCTAA 10020
 AGAGCTTCTT TTATCAGACT TTTCTTTGCT GAGTGAATTT CTCTTATCCT CCAAAGCAGC 10080
 CAAACGGGCT TGTGCCTCCT GTTGATTCAA GGCCACTTGC TCGGACTCCA GTTTCGATAG 10140
 GGCTAATTTT CTTTCTAAAT CACTAATCAG ACTAGTCAAG TCCATCAAAC TGCCTTGGTC 10200
 TTTGGCCATT TCAGCCTGTA AATCTTGGCG TTGCTTTTTA AGAGTTTGAT TTTCTTCTTC 10260
 TAATTTTTC ACGTTTGGT AATAACTCAT CAAGAGTTCT TGAACCTGAG TCAACTCTTC 10320
 TTCTGTCGAC TCTAGTTCAG CCTTATTTTC CTTGATTTGA GCAACCAGAA CATCTAAATA 10380
 AATAGCCTTA CGTTGTCCTT CCAAGTCTAA AACTTACGG GCATTCCTCAG CTGCTTCTC 10440

875

AAGAGGCTTG	ATTTGATTAT	CCAACCTCGTA	GATAATGTCC	TCTAAGCGGT	CCAGATTATC	10500
CTGAGTTTGC	TGCAGTTTAC	TCTCGGTTTC	TTTTCTGCGA	GTCTTGTATT	TTAAAACCTCC	10560
AGCAGCTTCT	TCAAAAATAG	CTCGTCGTTC	CTCAGGCTTG	GAATTA AAAA	TCTCCTCAAC	10620
CTTCCCTTGG	GAAATAATAG	AGAAGGAATC	TCGTCCCAAT	CCAGTATCCA	AGAAGAGGTC	10680
ATGAATATCA	CGCAGACGGA	CTTTCTTGCC	GTCAATCTTG	TATTCGCTAT	CTCCACTACG	10740
ATAGACATGG	CGTTCCACCC	TGATTTCTTG	ACCTGCATCC	TTGATAAATC	CGTCATGATT	10800
ATCCAGAGTC	ACAACCTACAG	AAGCATAATT	GAGCGGTTTG	CGACTTTCGG	TTCCAGCAAA	10860
GATGATATCC	GGCATCTTGC	CCCCACGGAG	ACTCTTGACA	CTAGACTCCC	CCAAAGCCCA	10920
ACGCAGACTT	TCTGTAATAT	TGGACTTTCC	AGATCCATTG	GGTCCAACAA	CTGCCGTCAC	10980
ACCTTGGTCA	AAAACGACCT	TGGTCTTATC	AGCAAAAGAC	TTGAACCCCT	GAATTTTCGAT	11040
TTCTTTTAAA	TACATGAATC	CAGCCCCCTC	TCAACGGCAT	TTTTTGGCAGC	TTCTTGCTCT	11100
GCTAATTTCT	TAGAACGACC	TTGGCCTTGA	CCGATGTCTT	TACCTTCAAC	AAGAACTTCT	11160
ACATCAAAAA	CCTTATCGTG	AGCAGGCCCT	GTTTCAGAAA	TCACCTGATA	ACGAATAGCC	11220
ACATCACCAT	TGACCTGAAG	CAACTCTTGG	AGATGGGTTT	TATAGTCTGT	AATCATCTCA	11280
AACTCGCCTG	CTTCAACCTT	AGGAATCATG	ACTTGATAGA	TAAATTCCTT	GACCTTGGCC	11340
ACATCCTTAT	CCAAAAGAAG	GGCACCAAGA	AAGGCTTCAA	AGGCATCACC	AAGAATGGTG	11400
TCACGATTGC	GACCACCTGA	TTTTTCTTCC	CCTTTACCCA	ACTTGATAAA	CTGGTCAAAC	11460
TGGCAATCAC	CGCAAAACC	AGCTAAACTC	TCCTCACGGA	CAATCATAGC	ACGGAGTTTT	11520
GATAGGTCAC	CTTCAGGCTT	TTTAGGATAT	TTTTTATATA	GATATTCTGA	AATCAATAAC	11580
TGTAGAACAG	CGTCTCTTAA	AAATTCCAAG	CGTTCATTGT	GTGAAATTTT	TAAGAGGCGG	11640
TGCTCATTTG	CATAACTCGT	ATGAGTAAAG	GCAGTTTCCA	GTAACCTTTT	GTCTGCAAAT	11700
TCGATTGCAA	AATGATCTTT	TAGTACAGTT	TGTAATCTTT	TCATACCAAC	CTCTTTCTAA	11760
CTGATAATAG	TCCTTTTTAT	TATATCAAAA	AAAGCCCCTT	GAGTCACTCT	AAAACGGGAC	11820
TGGAAAGCAT	TTGGGAATTC	TTTAGACAGA	GATTCTCAGT	TTTAGCGGCA	AATTTGGGTC	11880
AGGATAAAGA	AAAAAGCCCT	ATTAAAGGCT	TTTTAGGATG	TTTACATCCA	CCCTGAGGGA	11940
ATCGAACCCC	CATCTCAAGA	ACCGGAATCT	TACGTGATAT	CCATTACACT	AAGGGTGGAA	12000
ACTTGTTTTA	TTATAACAGA	AATTTGCTCT	AATAACAAGT	TTTTTGGTCA	AAGACCCCGT	12060
CTTAGTGGGA	AGCATCCCCA	TTCCAGATGG	AGTTTTTCAC	GATCACATAA	TCAACGTGTT	12120
TAAGGTCAGC	AACCTGACGT	CCACCTGCAT	AAGAAATAGC	ACTTTGAAGG	TCTTGTTCOA	12180

876

TCTCAGTTAA AGTGTCTTGC AGATGACCTT TAGCAGGAAG CAAGATACGT TTGCCTTCCA 12240
 CATTTTTGTA AGCACCTTTT TGATATTGTG AGGCTGAACC ATAATATTCT TTGAACTGTT 12300
 CACCATCGAC TTCAATCGTT TTCCCTGGAC TTTCAATGTG TCCTGCAAAG AGGGAACCAA 12360
 TCATGATCAT GCTAGCACCG AAGCGGATAG ACTTAGCAAT ATCACCGTGA GTACGAATTC 12420
 CTCCATCAGC GATAATCGGT TTACGCGCAG CCTTGGCACA CCAGCGTAGA GCAGCCAACT 12480
 GCCAACCACC TGTACCAAAA CCAGTCTTAA CCTTGGTGAT ACAAACCTTA CCAGGACCGA 12540
 TTCCGACCTT AGTAGCATCC GCACCAGCAT TTTCCAATTC ACGCACAGCT TCTGGTGTTC 12600
 CCACATTTC AGCAATGACA AAGGTATCTG GCAATTCTTT CTTGATGTGT TGAATCATAG 12660
 AAATCACGCT ATCCGCATGA CCATGAGCAA TATCAATAGT GATATACTCA GGAGTATCAG 12720
 CCTTGAGCTG GCTAACAAAA TCATACTCAT AATCCTTAAC ACCGACAGAG ATAGAAGCAA 12780
 TGAGCCCTTG ATTGTGCATT CGTTAATAA AAGGAATGCG TCCTGCCTCA TCAAAACGGT 12840
 GCATAATGTA GAAGTAACCA CCTTTAGCCA GTTGCTCTGC TACATTTTCA TCCAAAATCG 12900
 TCTGCATATT CGTGGCACA ACAGGTAGTT TAAAGGTGTG ATTTCCATAA GTGACACTTG 12960
 TATCCGCTTC TGCACGGCTT TTAATGACAC ATTTATTTGG AATCAATTGA ATATCTTCGT 13020
 AATCAAAAAT TGGAAATTC TTTAACATAT CGATGTCTCG TTTCTTTTGT AATGACCTAC 13080
 CTATGCTCTT GCATCACTAC GCCTTTTCCG ACGTTTCCTG G 13121

(2) INFORMATION FOR SEQ ID NO: 127:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 9578 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 127:

CCGAATGCAA TGTTTACGGT TGAAGTTGAA AATGGACATC AGATTTTAGC AACAGTTTCT 60
 GGTAATAATC GTAAAACTA TATTCGTATT TTAGCGGGAG ATCGTGTAC TGTCGAAATG 120
 AGTCCATATG ACTTGACAG TGGACGTATC ACTTACCGCT TTAAATAATC GAAAAACTTG 180
 GAGGGATAAG AAATGAAAGT AAGACCATCG GTCAAACCAA TTTGCGAATA CTGTAAAGTT 240
 ATTCGTCGTA ATGGTCGTGT TATGGTAATT TGCCCAGCAA ATCCAAAACA CAAACAACGT 300
 CAAGGATAAG ATAGAAAGGA GAAAACATGG CTCGTATTGC TGGAGTTGAT ATTCCAAATG 360
 ACAAACCGGT AGTAATCTCA TTGACTTATG TTTATGGTAT CGGACTTGCA ACATCTAAGA 420
 AAATTTTGGC TGCTGCTGGA ATCTCAGAAG ATGTTCTGTG ACGTGATCTT ACATCAGATC 480

877

AAGAAGATGC TATCCGTCGT GAAGTGGATG CAATCAAAGT TGAAGGTGAC CTTGTCGCTG	540
AAGTAAACTT GAACATCAAA CGTTTGATGG AAATCGGTTC ATACCGTGGT ATCCGTCACC	600
GTCGTGGACT TCCTGTCCGT GGACAAAACA CTA AAAACA CGCCCGCACT CGTAAAGGTA	660
AAGCTGTTGC GATTGCTGGT AAGAAAAAAT AATATAGGAG GTAAAAGTCT TGGCTAAACC	720
AACACGTAAA CGTCGTGTGA AAAAGAATAT CGAATCTGGT ATTGCTCATA TTCACGCTAC	780
ATTTAATAAC ACTATTGTTA TGATTACTGA TGTGCATGGT AATGCAATG CTTGGTCATC	840
AGCTGGTGCT CTTGGTTTCA AAGTTCTCG TAAATCTACA CCATTGCTG CTCAAATGGC	900
TTCTGAAGCT GCTGCTAAAT CTGCACAAGA ACACGGTCTT AAATCAGTTG AAGTTACTGT	960
AAAAGGTCCA GGTTCCTGGT GTGAGTCAGC TATTCGTGCG CTTGCTGCCG CTGGTCTTGA	1020
AGTAACAGCA ATTCGTGATG TGA CTCCAGT GCCACACAAT GGTGCTCGTC CTC AAAACG	1080
TCGCCGTGTA TAATCATCGC ATTACACTGC TTTTCGTTTA AGAGGGAGTA ACTAAATGAT	1140
CGAGTTTGAA AAACCAAATA TAACAAAAT TGATGAAAAT AAAGATTATG GCAAGTTTGT	1200
AATCGAACCA CTTGAACGTG GCTACGGTAC AACTCTTGGT AACTCTCTC GTCGTGTACT	1260
TCTAGCTTCT CTACCAGGAG CAGCTGTGAC ATCTATCAAC ATTGATGGTG TGTACATGA	1320
GTTTGACACA GTTCCAGGTG TTCGTGAAGA CGTGATGCAA ATCATTCTGA ACATTAAAGG	1380
AATTGCAGTG AAATCGTACG TTGAAGACGA AAAAATCATC GAACTGGATG TTGAAGGTCC	1440
TGCTGAAGTA ACAGCTGGTG ACATTTGAC AGATAGCGAT ATTGAAATG TAAATCCAGA	1500
TCATTATCTC TTTACAATCG GTGAAGGTTT TTTCTAAAA GCGACTATGA CTGTTAACAG	1560
TGGTCGTGGA TATGTACCTG CTGATGAAAA TAAAAGGAT AATGCACCAG TTGGAACACT	1620
TGCTGTAGAT TCTATTTATA CACCAGTTAC AAAAGTCAAC TATCAAGTGG AACCTGCTCG	1680
TGTAGGTAGC AATGATGGTT TCGACAAATT AACCCTTGAA ATCTTGACAA ATGGAACAAT	1740
TATTCCAGAA GATGCTTTAG GGCTTTCAGC ACGTATTTG ACAGAACATC TTGATTTGTT	1800
TACAAATCTT ACTGAGATTG CTAAGTCAAC TGAAGTGATG AAAGAAGCTG ATACTGAATC	1860
TGACGACCGT ATTTTAGATC GTACGATTGA GAACTGGAC TTGTCTGTGC GTTCATACAA	1920
CTGTTTAAAA CGTGCCGGTA TCAATACTGT GCATGATTTG ACAGAAAAAT CTGAAGCAGA	1980
GATGATGAAA GTACGAAATC TTGGACGCAA GAGTTTGAA GAAGTGAAAC TCAAACCTCAT	2040
TGATTTGGGT CTTGGATTAA AAGATAAATA AAGGAGGAAT ACATGGCTTA CCGTAAACTA	2100
GGACGCACTA GCTCACAACG TAAAGCAATG CTTGCGGATT TGACAACCTGA CCTTTTGATC	2160
AACGAATCAA TCGTGACAAC TGAAGCTCGT GCTAAAAGAAA TCCGTAAAAC TGTGAAAAA	2220

878

ATGATTACTC TAGGTAAACG TGGTGATTTG CATGCACGTC GTCAAGCAGC TGCTTTCGTA 2280
 CGTAATGAAA TCGCATCTGA AAACATGAT GAAGCAACTG ATAAGTACAC TTCTACTACA 2340
 GCACTTCAAA AATTGTTCTC AGAAATCGCA CCTCGTTATG CTGAACGTAA CGGTGGATAC 2400
 ACTCGTATCC TTA AAACTGA ATCACGTCGT GGTGATGCAG CGCCAATGGC GATCATCGAA 2460
 TTAGTATAAA ATCATCAATT TTGTGAGTG TTATGATGAT GGAGTCTTGT GCTCTTAGTC 2520
 TAGCTCTGGT CTACCGCTAG GATTTCGGTC CTAGCGGAA CACTCATCAT AAGTTGGGAT 2580
 AGTAGACGCT TGTTFACGAA ATTGTTTTTT TCTTAAGAAC AACTTCGTAA GCAGGCGTTT 2640
 TTGAGTATTT TCGTTAGAAT TATGCTATAC TATTTGAAAA GAATCCTGTT TAATGTTAAG 2700
 GTTCTTATTT TTAAGAAGAA TTGGAGTTTA CTTATGAAAG CCATTATAAC TGTGTGTTGGT 2760
 AAAGATAAAT CTGGAATTGT TGCAGGTGTT TCTGGTAAAA TTGCAGAATT AGGATTGAAT 2820
 ATTGACGATA TCTCTCAAAC TGTCTTGGAT GAATATTTTA CGATGATGGC TGTGTATCT 2880
 AGTGATGAAA AGCAAGATTT TACCTATCTT CGTAATGAAT TTGAAGCTTT TGGGCAAAC 2940
 TTGAATGTAA AAATCAATAT TCAGAGTGCA GCGATTTTCG AAGCTATGTA TAATATCTAG 3000
 GAGGTCATCA TGGATATTAG ACAAGTTACT GAAACCATCG CCATGATTGA GGAGCAAAC 3060
 TTCGATATTA GAACCATTAC CATGGGATT TCTCTTTTGG ACTGTATCGA TCCAGATATC 3120
 AATCGTGCTG CGGAGAAAAT CTATCAAAAA ATTACGACAA AGGCGGCTAA TTTAGTAGCT 3180
 GTTGGTGATG AAATGCGGC TGAGTTGGGA ATTCCTATCG TTAATAAGCG TGTATCGGTG 3240
 ACACCTATTT CTCTGATTGG GGCAGCGACA GATGCGACGG ACTACGTGGT TCTGGCAAAA 3300
 GCGCTTGATA AGGCTGCGAA AGAGATTGGT GTGGACTTTA TTGGTGGTTT TTCTGCCTTA 3360
 GTACAAAAAG GTTATCAAAA GGGAGATGAG ATTCTCATCA ATTCCATTCC TCGCGCTTTG 3420
 GCTGAGACGG ATAAGGTCTG CTCGTCAGTC AATATCGGCT CAACCAAGTC TGGTATTAAT 3480
 ATGACGGCTG TGGCAGATAT GGGACGAATT ATCAAGGAAA CAGCAAATCT TTCAGATATG 3540
 GGAGTGCCCA AGTTGGTTGT ATTCGCTAAT GCTGTTGAGG ACAATCCATT TATGGCGGGT 3600
 GCCTTTCATG GTGTGGGGA AGCAGATGTT ATCATCAATG TCGGAGTTTC TGGTCCTGGT 3660
 GTTGTGAAAC GTGCTTTGGA AAAAGTTCGT GGACAGAGCT TTGATGTAGT AGCCGAAACA 3720
 GTTAAGAAAA CTGCCTTTAA AATCACTCGT ATCGGTCAAT TGGTTGGTCA AATGGCCAGT 3780
 GAGAGACTGG GTGTGGAGTT TGGTATTGTG GACTTGAGTT TGGCACCAAC CCCTGCGGTT 3840
 GGAGACTCTG TGGCACGTGT CCTTGAGGAA ATGGGGCTAG AAACAGTTGG CACGCATGGA 3900
 ACGACGGCTG CCTTGGCCCT CTTGAACGAC CAAGTTAAAA AGGGTGGAGT GATGGCCTGC 3960
 AACCAAGTCG GTGGTTTATC TGGTGCCTTT ATCCCTGTTT CTGAGGATGA AGGAATGATT 4020

GCTGCAGTGC	AAAATGGCTC	TCTTAATTTA	GAAAACTAG	AAGCTATGAC	GGCTATCTGT	4080
TCTGTTGGAT	TGGATATGAT	TGCCATCCCA	GAAGATACGC	CTGCTGAAAC	TATTGCGGCT	4140
ATGATTGCGG	ATGAAGCAGC	AATCGGTGTT	ATCAACATGA	AAACAACAGC	TGTTTCGTATC	4200
ATTCCCAAAG	GAAAAGAAGG	CGATATGATT	GAGTTTGGTG	GTCTATTAGG	AACTGCACCC	4260
GTTATGAAGG	TTAATGGGGC	TTCGTCTGTC	GACTTCATCT	CTCGCGGTGG	ACAAATCCCA	4320
GCACCAATTC	ATAGTTTAA	AAATTAAGAA	AATAGGAGAA	ATTTTAAGTT	CTATTTAAGA	4380
TTAGACGTGT	ATACTATAAT	CATTAAATA	AGACCTCCTA	ATATTATTTG	AAACAGATAA	4440
CACTGAATTA	GTTTGAATTT	GATTTTCATC	TAATATCTTT	ATTTAATGAA	CTCCTAAACT	4500
TTTTCATAAT	AATCTCCTTC	AAAAGTCGCC	TGTATGGGTG	GCTTTTATTT	TATCATTCAT	4560
GATATAATAG	AAGCAAACGG	AGGACGGAAA	ATGGTAAAAG	TACGATTGTA	TTTGGTACGT	4620
CATGGCAAGA	CCATGTTTAA	CACGATTGGT	CGCGCGCAAG	GTTGGAGCGA	TACTCCCTTA	4680
ACTGCTGAAG	GTGAACGAGG	GATTCAAGAG	TTAGGAATCG	GTTTGCAGGA	ATCTGATCTA	4740
CAGTTTGAGC	GTGCTTATTC	GAGTGATTCT	GGTCGTACCA	TTCAGACCAT	GGGAATTATC	4800
CTTGAAGAAC	TTGGCTTGCA	GGGGGAAATC	CCTTATCGCA	TGGACAAGCG	TATCAGAGAA	4860
TGGTGTTCG	GTAGTTTGA	TGGAGCCTAT	GATGGCGATC	TTTTCATGGG	CATTATTCCT	4920
CGTATCTTTA	ATGTGGACCA	CGTTCACCAA	TTGTCTTATG	CTGAACTGGC	TGAGGGCTTG	4980
GTAGAGGTCG	ATACAGCTGG	TTGGGCTGAA	GGCTGGGAAA	AACTCAGTGG	CCGAATCAAG	5040
GAAGGCTTTG	AAATGATTGC	AAAAGAAATG	GAAGATCAAG	GTGGAGGTAA	CGCCCTTGTT	5100
GTCAGCCATG	GAATGACTAT	TGGAACCATT	GTTTATCTGA	TTAATGGCAT	GCATCCGCAT	5160
GGTCTGGATA	ATGGTAGCGT	GACAATCCTT	GAATATGAGG	ACGGCCAGTT	TAGGGTTGAA	5220
GTTGTGCGTG	ACCGTAGTTA	CCGAGAGCTA	GGACGTGAGA	AGATGGAAGA	AGGCTCTATT	5280
TAATCAGTCT	AGACTTGCTT	GCCATGAGCT	AGGGATTTGA	TAAGAATATC	AAGATAAGAA	5340
AAAACAGCCG	AGGGCACTCC	TTTCGGCTGT	TTTTGATGTG	GAAAACTAAA	GTGTAATGCT	5400
ATTGCTTTTA	GAGATTTTCA	TAAACAAGAG	CAAGGAACCT	ACTGTTAGAA	CAGTCAGGAT	5460
AGTTGACAAG	GTTCGGGCTA	CACCGTAATT	TCCTCTGAGA	ACCTCTGTAT	AAATAGCTAC	5520
AGTCATTGTT	CTTGTTTTGA	CATTGTAGAG	GAGGATAGAA	GTAGAGAGTT	TTGAAATCAT	5580
TGTGACTCAA	GATAAGATGG	CTCCAGAAAT	GATACCAGAT	AGCATCATTG	GAGTTGTAAT	5640
CTTAGCAAAG	GTATTGAGAC	GACTACTTCC	TAAGCTTTCA	GCAGCTTCTT	CAATACTTGG	5700
TGCTATTTGT	TGTAAGCTAG	CAACAGATGA	GCGAATAGTA	TAAGGTAATC	TTCTGGCAGA	5760

TAGAGACATA ATCAAGATGA AAGCAGTCCC TGTAATCATA AGAAATCCAC TTCCAAATAG 5820
 ACCAGTATTG AAGGAAGAAA TGAAGGCAAT CCCTAGAACG GTTCCTGGTA CAATATAAGG 5880
 TACCATACTG AGGCTGTCAA TTAAGTTTGT AAACAAATTC CGTTTTCTAA CGGCTAGGTA 5940
 GGAGATAAAT GTCGCAAATA GAACAAC TAG AACTAAGGCA ATCAAAGGGA TACGAATGGT 6000
 ATTGAAAATA GCAGATCCCA TACGATGGAA AGCTACCTTG TAACTGTTTG GAGAATAACC 6060
 TTTAACAGAT ACCATACCTG ATGTTTTTAG GAAAGAGGTA TAAATTAAGT AGATTTGAGG 6120
 TAAAACAGAG ATAAAGATAA TTCCGTAGAC TGTTGCATAA ATGGCAGCCA TTTTTCCTTT 6180
 TGTAGTTTTT TTAGGCTCAA TTGGATGGAG CAGATTCATG CTGAAACTGT AGCGGTTTTGC 6240
 AATGTGTTTT TGGATAAGGA AAATTGCCAA GGCAATGATA ATCGCCATAA TTGCAAAAGC 6300
 AGAATTTCCCT CCAACCTCGC TAATAAATG GGTATAAATC AGGACAGGGA AAGTCCGATA 6360
 CCCTTCGCCA ATCAACATAG GCGTTCCAAA GTCTGAGAAT GCTCTCATAA ATACAAGCAA 6420
 GGAGCTGCTA GTAAGGTTGG AACTAGGAGA GGTAACAACA CCGTTACGAT AGGTTTAAAT 6480
 CCGAAGGACC CCATGCTTTC AGCTGCTTCA AGTAGAGAAT TGTCAATACT GTTCATTGTT 6540
 CCAGCAACAT ATAGAAATAC CAGTGGGAAT AGTTGCAGTG TAAAGACAAG TACAATTCCCT 6600
 TTGAATCAAT AAATATCGAT AGCTGGAAGA TAAAGGGCAT TTGTCAAAAA TTTAGTGATG 6660
 ACCTCATTTC GTCCTAGCAA GAGAACCAG GAGTAGGCTC CTACGAAAGG AGCTGACATG 6720
 GAAGCAATGA TAATCAATAT TTGTAGAAAT TTCTTCCCCT TGAAGTCATA CATAGAGAAG 6780
 AGATAAGCTA ATAGGGTTCC TACAAC TAAG GAAGTGATAG TAGCGGTAAT GGAAACCTTG 6840
 AAAGTGTGTA CTAGTGTCTC AGAGTAGTAG GCTTTACTAA AGAAAGTGAC AAAATTAGCT 6900
 AGTGAGAATT GTCCTCATG TATAAGTGCT TGCTTGAGCA CGGTAACGAT AGGATAAACG 6960
 AGAAAGATAG GATAGGTAAG AAAGAGGAAG AAAGAGGAAA CTGTCCAAAT ATTTAGTTTT 7020
 TTACGTCCA TGGTTGACTC CTTTTATCAG GTTTTGGGAA CCATCTGCAG AAAAGATGTT 7080
 TAATTTTTCG GTATTGATTC GTAGACGAAT ACGATTGCCT TTTTGTAGAT CTCTTCAAA 7140
 AGTTGATTCT TCACTA ACTT GAATTTTGA GGCAAAACCT GTCTCAATGA AATAATCCGT 7200
 ATTTAGTCCA AGATAGACGC TATCTCTAAT AGTTCCTTCA ATATCTCCAG ATTCATCTTT 7260
 GATAAACTCT TCGGGACGAA TGCTTACATG AATAGCTTGC TCCTCAACCT GATCAAGAGC 7320
 TGGCATTCGA AGGGCATAGC CATCTGAAAA GACGATATAA GCGCCGTCGC TCCGTTTTTC 7380
 AAGATTGGCA GGGATAATAT TTGTGCGTCC GATAAAGGTT GCCACAACT CATTAGCTGG 7440
 TTTATGATAG AGTTCTTTTG GTCGGCCGAT TTGTGGATC ACCCATCTT TCATAACAGC 7500
 AATTTGGTCT GAAATAGCCA TGGCTTCTTC TTGGTCTGTT GTTACATAAA CAGTTGTAAT 7560

881

TCCCCTTCG	TGTTGGATTT	CTCGGATGGC	TTGACGCATA	TCCAAGCGAA	GTTTGGCCTC	7620
CAGATTACTA	AGTGGCTCGT	CCATGAGGAG	AACACTTGGA	TTAACCGCTA	AGGCGCATGC	7680
CAAGGTGACA	CGTTGTTGTT	GTCCACCACT	GAGTTTATCG	GGCTTTCGAT	CCGCATATTG	7740
AGCAATTTGC	ATGAGTTCAA	GATACTTGTT	GGTCTGTGTA	ATCAATTCTT	CTTTTGGAAC	7800
CTTCTTTTGC	ATAAGACCAA	AAGCAACGTT	GTCTCGGACA	GTCAAATGTG	GGAAAATAGC	7860
GTAGTTTTGG	AAAACCATCC	CGATATTGCG	TTTGCTGGGT	TCCATATTAT	TGATTTTTGT	7920
ATCATCGAAG	TAAATTTCTC	CACCTTCGAT	ACTGTTGAAA	CCTGCAATCA	TACGAAGAAG	7980
GGTCGTTTTC	CCACATCTG	AAGCTCCAAG	AAGGGTAAAG	AGACTTCCTT	TTGGAATTGT	8040
AATGTTCAAA	TTCTCAATAA	CAGGGACATC	GTGGTAGATT	TTTTTGGCGT	TAATAATTTT	8100
GATCTCACTC	ATAGTGAACC	TCTTTTACTG	TTTAGATTGG	ATATCTGTAA	AGACTTCGTT	8160
GTATTTCTTA	ACGATATCTG	ATTTATTCTT	GATGACATAA	TCATAATCTT	CAGTGAGTGT	8220
TTTGATTTTG	TCAATTGTTT	TCATGTTTTC	GCTTGTTTTA	GCATTTTAC	GAACAGGACG	8280
GTTAGTAGTG	GTTGTACCAA	GTGTATCTTG	TACTTCTTGA	GAGATAATAA	AATCGATAAA	8340
TTTCTTGCCA	TTTTCCATAT	TTTTAGATTT	TTTAACGATA	GCAGCACTAG	CAGGTAGGAA	8400
GACGGTCCCT	TCTTTTGGAT	AGACTACCTT	AATGTTAGCT	CCGTCATTTA	AGAGTTTAAAC	8460
TGCTGGATCT	TCATAAGAGA	GACCAACAGC	CATTTCTCCA	TCAGCGACTA	CTTTATAGAC	8520
ACTAGATGAA	CTTGAACCGA	TTTTACCATC	AATAAGTGTG	AAAAGATCTT	TTACATAAGA	8580
CCAAGCCTTA	TCATCTTTGT	AACCACCTTG	AGCTTGTAGC	ATATTTGTTA	ATTGAGCAAA	8640
GGCGCTAGAA	GAGTTTGCTG	GGTCAGCAGT	TGCGATTTTT	CCTTTTAGTT	CAGGTTTGAA	8700
AAGATCGTTA	TATCCTTCGA	TGTTTATGCC	TTTAGTTAAA	TCAGGGTTGA	CGATTAATAAC	8760
ACTACCATCT	AGTGTATAAG	GAGTAGAGTA	GCCAGTTGTG	TTTTGATATT	CTTTGATAAC	8820
ATTATCATTT	TCTTTTGAAG	TATAGTTTTC	AAAGAGTTCT	CCGTGGGTAG	TATATTGTGT	8880
ATAAGAACCA	CAAAGATAA	CATCAGCTAC	AGGAACTTCT	TTTTCTGACT	CTAGTTTTTTT	8940
GAAAAGTTCT	CCAGTACCAG	CTTGAATCAG	TTCTACTTTG	ATACCATATT	TTTCTTCAAA	9000
GGCAGGAATA	GTTGCTCCAA	TTAAGCCCTC	TGAGTTTGGT	GAATAAACGA	CTAGCGAACC	9060
GCCGTCTCCT	TTATCAGATG	AACTGTCATC	GGCAGATTCA	TTAGAAGAAC	AAGCAGCATA	9120
ATACATCCAT	TTCTTTTTCA	TGATGGATAC	CTCCGTGTG	TTATTTAAGT	TTATTTTAAA	9180
ACAATGTAAG	CGTTTTTAAA	ACATACAATT	CTATTCTATA	GTGTATTGAA	TCTATAACAG	9240
TACACTTTGA	CTGCTAAAAT	ATTTCTATAA	ATTAATTGTA	CTTTCCTGAT	AGAGATGTTC	9300

882

ACATCTTATT TCAATTCACT ATATTAGAGT AAAATTCTCT ACAAAAAGAA GAATAGCCTA 9360
 TTTTACTATT CTTCTGAGTG ATTTCAATTC CTTTGGGGAA ATATGGAGAT ACTTTTTTAAA 9420
 TCCTGACAAA TGGTTGTTTC TTTTTCTAAA TCGGTGATAC TGTATCGGAG AATGCGCGTG 9480
 AGGTCACAAA GGCTGCGATA GAGCTTCTAT GGAGAATTC TTTTGGGAGA GATTTTTTAA 9540
 AGGAATGAGA CATCCGCTAC CTCCTTGAA GGTTTTTG 9578

(2) INFORMATION FOR SEQ ID NO: 128:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 13440 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 128:

CGGGCTGTTG TGACGATTCT TATTTCTATC TGTGTTATCT TTTTGGGAAC TATTTTGGGT 60
 GTTGTCTTGG CTTTTGGGCA ACGTTCAAAG TTAAACCGC TTGTTTGGTT GGCCAACTTG 120
 TACGTTTGGG TTTTCCGTGG GACACCGATG ATGGTTCAA TTATGATTGC CTTTGCTCTT 180
 ATGCATATCA ATGCTCCGAC TATTCAGATT GGAATTTTAG GTGTTGATTT TTCGCGCTCG 240
 ATTCCAGGGA TTTTGATTAT CTCTATGAAT AGTGGTGCTT ATGTTTCGGA GACTGTTTCGT 300
 GCCGGAATCA ATGCGGTTC AAAAGGTCAG CTAGAAGCGG CTTATTCGCT AGGGATTCGT 360
 CCTAAAAATG CGATGCGTTA TGTGATTTTG CCACAAGCAG TCAAAAATAT CTTGCCAGCA 420
 TTGGGGAACG AATTTATCAC CATTATCAAG GACAGCTCCC TCTTATCAGC TATTGGGGTC 480
 ATGGAGTTGT GGAATGGGGC TACAACAGTT TCTACAACAA CCTATCTACC TTAAACACCA 540
 CTTTTATTG CAGCATTTTA CTACTTGATT ATGACCTCTA TTCTGACAGT AGCCTTGAAA 600
 GCTTTTGAAA AACATATGGG ACAAGGAGAT AAGAAATAAT GACAGAAACC TTGATAAAAA 660
 TTGAAAATTT ACATAAATCC TTTGGAAAGA ATGAAGTATT GAAGGGCATC AACCTCGAGA 720
 TTAAAAGAGG AGAAGTTGTC GTTATCATCG GTCCTTCAGG GAGCGGGAAA TCTACCTTGC 780
 TTCGCTCTAT GAATTTGTTG GAAGAAGCAA CCAAGGGGAA GGTATCTTT GAGGGAGTCG 840
 ATATTACGGA CAAGAAGAAT GACCTGTTTG CCATGCGTGA GAAGATGGGC ATGGTTTTTC 900
 AACAAATCAA TCTCTTCCT AATATGACTG TGATGGAAAA TATCACCTTG TCCCCTATCA 960
 AGACCAAAGG TGACAGTAAG GCCGTTCAG AGAAAAGAGC TCAGGAAGTT TTGGAAAAAG 1020
 TTGGTTTGCC AGATAAGGCA GACGCTTATC CACAGAGTTT GTCAGGTGGC CAGCAACAGC 1080
 GGATTGCCAT CGCGCGTGGG TTGGCTATGG AACCAGATGT TTTGCTCTTT GACGAGCCAA 1140

CTTCAGCCCT	AGATCCTGAG	ATGGTTGGAG	AAGTTCCTGGC	TGTTATGCAA	GATCTAGCCA	1200
AGTCAGGAAT	GACCATGGTT	ATCGTAACAC	ATGAGATGGG	ATTTGCCCGT	GAGGTGGCAG	1260
ATCGTGTTCAT	CTTTATGGCA	GACGGTGTGG	TTGTTGAAGA	CGGAACACCT	GAGCAGATTT	1320
TTGAACAAAC	CCAAGGACAA	AGGACTAAAG	ACTTCTTGAG	TAAGGTTTFA	TAAGTTAGCT	1380
TTGTTTAGCT	ATTTGTAGCC	AGCTTTAAAC	GTAAAGAGA	AGATTAGTGA	AAAGCTCAAC	1440
CAGAGCTTTT	TCTTATAGTT	TAAAGCTATA	GGATTGCCCTA	GGAAAGAAGT	GTTAGAGCTA	1500
CATTGTATTT	TTTGGTATAA	TTAAAGATAT	TTGTAAGAAA	AGAGAAGTGA	TATGACACAG	1560
ATTATTGATG	GGAAAGCTTT	AGCGGCCAAA	TTGCAGGGGC	AGTTGGCTGA	AAAGACTGCA	1620
AAATTAAAGG	AAGAAACAGG	TCTAGTGCCT	GGTTTGGTAG	TGATTTTGGT	TGGGGACAAT	1680
CCAGCCAGCC	AAGTCTACGT	TCGCAACAAG	GAGAGGTCAG	CCCTTGCGGC	TGGTTTCCGT	1740
AGCGAAGTAG	TACGGGTTCC	AGAGACCATT	ACTCAAGAGG	AATTGTTAGA	CCTGATTGCT	1800
AAATACAATC	AGGATCCAGC	TTGGCATGGG	A'TTTTGGTTC	AGTTGCCATT	ACCAAAACAC	1860
ATTGATGAAG	AGCGGT'TCT	ATTGGCTATT	GACCCAGAAA	AGGATGTGGA	TGGTTTCCAT	1920
CCTCTAAACA	TGGGGCGTCT	TTGGTCTGGT	CATCCAGTCA	TGATTCCTTC	GACACCGGCA	1980
GGAAT'TATGG	AAATGT'TCCA	TGAATATGGG	ATTGACTTGG	AAGGTA'AAAA	TGCAGTCGTC	2040
ATCGGTTCGAT	CCAATATTGT	CGGAAAACCT	ATGGCCACAGC	TTCTTTTGGC	AAAGAATGCA	2100
ACAGTAACCT	TGACTCACTC	ACGTACTCAT	AATCTT'TCCA	AGGTGGCTGC	AAAAGCAGAT	2160
ATTCTGGTTG	TTGCAATCGG	TCGTGCCAAG	TTTGTGACTG	CTGACTTTGT	CAAACCAGGT	2220
GCGGTAGTCA	TTGACGTTGG	GATGAACCGC	GATGAAAATG	GTAAGCTCTG	TGGGGATGTT	2280
GATTATGAGG	CGGTTGCCCC	ACTTGCTAGC	CACATTACGC	CAGTCCCTGG	AGGTGTCCGGT	2340
CCTATGACCA	TFACTATGCT	GATGGAGCAA	ACCTATCAGG	CAGCACTTAG	GACATTGGAT	2400
AGAAAATAAG	ATAAAAATTT	TCTGAGGAAA	GTGTATTTTC	TATAGCTATA	TCTAAAATGA	2460
TAGAAATGAA	TATTAATTTT	TAGAAATAAG	TTTATAAAAAG	GAGGTTTGGC	CCTCCTTTTT	2520
GTTGTATAAT	GGAGTGAGGT	GATTAGATGA	TTTTAAAAAT	TTATAATGGG	GAATATAGTT	2580
TACAA'TGGGA	TGGAATATAC	TACTTAGCAC	TAATTGATTA	TCCAAATATT	CAAGAGTGGG	2640
AATTAGAAAA	AATTGCTAAA	TTTATAGCTT	ACGAAAAACT	TCATAAACGT	CAAACAAGTA	2700
TTGAGTGTGC	TGATTC'TTGT	TTAAAAAAG	AAATTTTAGA	TTACATCTGT	CAGCATCCCT	2760
TTCTGCCACC	ATTTACTCCT	ACAGATAAAA	GAGTAGCCTC	GACTTATGAC	CTACATAAGA	2820
GGTTAGTGAC	TTCAGACTAC	TGTAGTCATA	CTACGACTAT	AGATGCAGCG	ATTTCTATTT	2880

TTAAAACCTGG	TCGCTTTTTA	TCTGCTGTGA	AAGCCTTTGG	GCGAGATGCT	GAGGAGTTGG	2940
TTTTGGATAG	TCGAAATGCT	GCATCTGATC	CGATAGATTA	TTTTGACTAT	GTCATGTTAG	3000
GGTGGTCAAA	TACAAGTTCT	GGTTATCGAT	TGGCGATGGA	GCGTTTATTA	GGTCGAGCTC	3060
CTTCAGAGAA	AGAATTACAA	GACAAGTTTA	TTCCCTGGAGT	AAGTTTTTCAT	TTTATCTATA	3120
CAGATTTGAT	TAAAGTTCCT	GGTTATATTT	TTGATGGTTA	CCATGCTGTA	AAAATTAAGG	3180
ACATGCTTAA	TTTATTAAGT	GAGTTGTATA	TTTGCATTAT	TCCAACCTCAT	AATAAGAGCC	3240
AATTTGAAAA	TATTATTTCA	ACCAAAATAC	AAGATAGGGT	GTATTATCTT	GACTATGCTG	3300
GAGAAGACTT	AGAAGAGTGG	ACTAAGAAAAG	TCTATCAAGT	TGTTTTAAAA	CAATCAGATA	3360
AAGGATAGTT	GAGGAAAAAA	CGATGAAAGT	GATTGATCAA	ACCTTACTAG	AAAAAGTCAT	3420
TATTGAACGT	TCTTGTAACA	GTCATAAAGG	AGACTACGGT	CGTCTGCTGT	TGCTTGGTGG	3480
GACTTATCCT	TATGGTGGTG	CCATCATCAT	GGCTGCTTTA	GCAGCTGTAA	AAAGCGGTGC	3540
AGGATTGGTA	ACCGTTGGAA	CGGACAGGGA	AAATATCCCT	GCTCTACACA	GCCATTTGCC	3600
TGAGGCTATG	GCCTTTTCTC	TGCAAGATCA	GTAATTGTTA	CAAGAGCAAT	TGGAGAAGGC	3660
AGAAGTTGTC	TTGCTGGGGC	CTGGTTTACG	AGACGATACG	TTTGGAGAAA	ATCTTGTAAG	3720
ACAGGTCTTT	GCTAGCTTAA	AAAAGAATCA	GATTTTGATT	GTAGATGGAG	GGGCCTTAAC	3780
CATCCTTGCT	AGGACAAGTT	TGTTGTTTCC	ATCTAACCAG	CTTATCTTAA	CTCCCCACCA	3840
AAAAGAATGG	GAAAAACTGT	CTGGTATTGC	TATTGAAAAG	CAAAACGAAG	GTACAACATC	3900
TAGTGCCCTG	ACTTCTTTCC	CTCAAGGAAC	AATTTTGTTA	GAGAAAGGTC	CAGCTACTCG	3960
TATTTGGCAA	GTTGGCCAGT	CTGATTATTA	CCAGTTAAAG	GTTGGCGGTC	CCTATCAGGC	4020
GACTGGTGGT	ATGGGTGATA	CACTGGCTGG	AATGATTGCA	GGATTTGCAG	GCCAATTTCCG	4080
ACAGGCCAGT	CTCTACGAAC	GTGTGGCAGT	AGCAACCCAT	CTTCATTTCAG	CCATAGCCCA	4140
AGAACTATCT	CAAGAAAATT	ATGTGGTCTT	GCCGACGGAA	ATTAGTAATT	GTCTTCCTAA	4200
AGTAATGAAA	AGATATGTCT	AAAATAGTTA	GACAAAAAAT	GTTGATAAAT	TGTATCATTAA	4260
TTCTTAATTC	ACAAAAAACG	AACGTTTAGT	ATTCTTCTTG	CTAAGAAACT	AAATTTGTTT	4320
GTTTTTTTAC	TCTTGTAAT	CTATTTTTGT	TAGAGTTGAT	TTGGTTTACA	TCCGTACTTA	4380
AATTGATTTG	TTAGAGCTCT	ACTTTTATTA	AAAAAATTCA	ATTTCAAGGA	TAAATAAGCA	4440
GTATTCTAAA	GGTACTTTTA	GATGAAATAA	AAGCCTTTAC	ATGGTATAAT	AGAGGTAGCT	4500
CTTTAATGGA	GGTGTGAG	TGGAAAATCT	GAAGAAAATG	GCAGGTATCA	CGGCTGCTGA	4560
ATTTATCAAG	GATGGGATGG	TTGTAGGGCT	AGGAACAGGT	TCTACTGCCT	ATTATTTTGT	4620
CGAAGAAATC	GGTCGTCCAA	TCAAGGAAGA	AGGCTTGCAG	ATTACAGCTG	TGACGACTTC	4680

TAGTGTGACC AGTAAACAGG CTGAAGGGCT CAATATCCCG CTCAAGTCTA TTGACCAAGT 4740
 AGACTTTGTC GATGTGACAG TCGACGGGGC GGATGAAGTG GATAGTCAGT TTAATGGAAT 4800
 CAAAGGCGGT GGTGGTGCCC TTCTCATGGA AAAGGTGGTC GCAACACCAT CAAAAGAATA 4860
 CATTTGGGTG GTGGATGAAA GCAAGCTGGT CGAAAACTA GGTGCTTTTA AATTGCCAGT 4920
 AGAAGTGGTT CAGTATGGTG CAGAGCAGGT CTTTCGTCAT TTTGAACGAG CTGGCTACAA 4980
 ACCAAGTTTC CGTGAAAAAG ACGGCCAACG TTTTGTGACC GATATGCAGA ATTTTATCAT 5040
 TGACCTCGCC TTGGATGTCA TTGAAAATCC AATTGCTTTT GGACAAGAAT TGGACCATGT 5100
 CGTTGGTGTG GTGGAGCATG GTTTATTCAA CCAAAATGGTG GATAAGGTAA TCGTTGCTGG 5160
 ACGAGATGGA GTTCAGATTT CAACTTCAAA AAAAGGAAAA TAGAAGGGGG CATAAGATGT 5220
 CTAATTTTAA TCGTATTCAT TTGGTGGTAC TGGATTCTGT AGGAATCGGT GCAGCACCAG 5280
 ATGCTAATAA CTTTGTCAAT GCAGGGGTTT CAGATGGAGC TTCTGACACA CTGGGACACA 5340
 TTTCAAAAAC AGTTGGTTTG AATGTCCCAA ACATGGCTAA AATAGGTCTT GGAAATATTC 5400
 CTCGTGAAAC TCCTCTTAAG ACTGTAGCAG CTGAAAGCAA TCCAACCTGA TATGCAACAA 5460
 AATTAGAGGA AGTATCTCTT GGTAAGGATA CTATGACTGG AACTGGGAA ATCATGGGAC 5520
 TCAACATTAC TGAGCCTTTC GATACTTCTT GGAACGGATT CCCAGAAGAA ATCCTGACAA 5580
 AAATCGAAGA ATTCTCAGGA CGCAAGGTTA TTCGTGAAGC CAACAAACCT TATTCAGGAA 5640
 CGGCTGTTAT CTATGATTTT GGACCACGTC AGATGGAAAC TGGAGAGTTG ATTATCTATA 5700
 CTTCAGCTGA CCCTGTTTTG CAGATTGCTG CCCACGAAGA CATTATTCCT TTGGATGAAT 5760
 TGTACCGTAT CTGTGAATAC GCTCGTTCGA TTACCCTTGA GCGTCCTGCC CTCTTGGTGTC 5820
 GCATCATTCG TCGCCCTTAT GTAGGTGAAC CAGGTAACCT CACTCGTACG GCAAACCGTC 5880
 GTGACTTGCC TGTATCTCCA TTTTTCCTAA CTGTTTTGGA TAAATTTGAAT GAGGCTGGTA 5940
 TCGATACTTA TGCTGTGGGT AAAATCAACG ATATCTTTAA CGGTGCTGGT ATCAACCATG 6000
 ACATGGGTCA CAACAAGTCA AATAGTCATG GAATTGATAC ACTATTGAAG ACTATGGGAC 6060
 TTGCTGAGTT TGAAAAAGGA TTCTCATTCA CAAACCTAGT TGACTTTGAT GCCCTTTACG 6120
 GCCATCGTCG TAATGCTCAC GGTTACCGTG ATTGCTTGCA TGAGTTTGAT GAACGCTTAC 6180
 CTGAAATTAT CGCAGCTATG AGAGAGAATG ACCTTCTCTT GATTACTGCG GACCATGGAA 6240
 ATGACCCAAC GTATGCAGGA ACGGATCACA CTCGGGAATA TATTCCATTG TTGGCCTATA 6300
 GCCCTGCCTT TAAAGGAAAT GGTCTCATTC CAGTAGGACA TTTTGCAGAT ATTTACGCGA 6360
 CTGTTGCCGA TAACTTTGGT GTGGAACTG CTATGATTGG GGAAAGTTTC TTAGATAAAT 6420

886

TGGTATAAGA	TGACGCGCTA	TGCTTTGCTG	GTGAGAGGTA	TCAATGTTGG	TGGTAAGAAT	6480
AAGGTCGTCA	TGGCGGAGCT	TCGTCAAGAA	TTGACAAACT	TGGGACTGGA	AAAGGTTGAG	6540
AGCTACATCA	ATAGTGGCAA	TATTTTCTTT	ACTTCGATAG	ATTCCAAAGC	CCAATTGGTT	6600
GAAAAGCTAG	AGACTTTCTT	TGCAGTCCAT	TATCCATTTA	TTCAGAGCTT	TTCTTTACTG	6660
AGTCTAGAGG	ACTTTGAGGC	GGAAC TTGAA	AATCTACCAG	CTTGGTGGAG	CAGAGACTTG	6720
GCACGAAAAG	ATTTTCTCTT	TTACACTGAG	GGTTTGGATG	TGGACCAAGT	CATCGCGACA	6780
GTTGAAAGTT	TAGAGCTGAA	AGATGAAGTG	CTTTATTTTG	GAAAAC TTGG	GATTTTCTGG	6840
GGGAAATTTT	CTGAAGAATC	CTATTCTAAG	ACTGCCTATC	ATAAGTACTT	GCTGAAGGTG	6900
CCTTTCTACC	GCCACATTAC	TATTCGTAAT	GCTAAACCTT	TTGACAAAAT	TGGTCAAATG	6960
CTAAAAAAT	AATAAAGGAG	ACACACAATG	ACATTTTAA	ACAAAATCCA	TGAAACTGCT	7020
ACTTTCCTGA	AAGAAAAGGG	AATTGCAGCC	CCTGAGTTCC	GTCTAATCCT	TGGATCAGGA	7080
CTTGGAGAAT	TGGCAGAAGA	AATCGAAAAT	CCAGTTGTAG	TAGACTATGC	TGAGATTCCA	7140
AACTGGGGCC	GTTCAACAGT	AGTCGGTCAT	GCTGGTAAAT	TGGTATATGG	TGAACTGGCA	7200
GGTCGCAAGG	TCTTGGCTCT	TCAAGGGCGT	TTCCATTTCT	ATGAAGGGAA	TCCTCTGGAA	7260
GTGGTGACTT	TCCCAGTTCG	TGTGATGAAA	GTTCTTGGAT	GTGAAGGTGT	TATTGTAACC	7320
AATGCAGCTG	GCGGTATCGG	ATTTGGTCCT	GGTACCTTGA	TGGCTATCTC	AGACCATATC	7380
AACATGACGG	GGCAAAATCC	ATTGATGGGT	GAAAAC TTGG	ATGACTTTGG	CCCACGTTTC	7440
CCAGATATGT	CTAGGGCCTA	CACACCAGAA	TACCGTGCCA	CTGCCCATGA	AGTGGCTAAA	7500
AAACTTAATA	TCAAGCTTGA	TGAAGGTGTC	TATATCGGAG	T TACTGGTCC	GACTTATGAA	7560
ACACCAGCAG	AAATTCGTTC	CTATAAGACA	CTGGGAGCAG	ATGCAGTTGG	TATGTCTACG	7620
GTTCTGAAG	TTATCGTGGC	AGCCCACTCT	GGCTTGAAAG	TTCTGGGAAT	TTCATGTATC	7680
ACTAACTTTG	CGGCCGGTTT	CCAAGAAGAA	CTCAATCACG	AAGAAGTTGT	AGAAGTGA CT	7740
GAACGTGTTA	AAGGTGATTT	CAAAGGCTTG	CTTAAAGCGA	TTCTTGCTGA	ATTGTAAGAA	7800
AAAAGATTTA	AAAGGGGGAG	TGCCTCTGTT	TTTT CAGGAT	TGACTGCCTA	TCCGGATTAA	7860
AGAAGAAACA	GAGGAATACT	ATGAGCTTCT	TCCTGCTCTT	ATAACTGAAA	GAAGCGGAAG	7920
AATAGGTATG	TCTGATCTGA	TAGCCAGCAT	TGTGAAAGAC	AAGATTCTAG	GATACTAGCA	7980
TTAGCTTCCT	AGCCAAGCAG	ACTAGTATGA	TAAGGAGAGA	TGAGAATGAA	TTGACTTTCT	8040
GAATTTCTCA	GTCTTATCAT	ATATAGCACA	ATGAGATTTT	GCTTGAGTCT	GCTTGTA AAT	8100
AAACGAAAAG	AAAGATAAGA	AATAATGAAA	ATTTGGTCAAC	GAATTATGCG	CTTTGGCATA	8160
AAAAATTAAG	TATCGGAGTT	GTATCTGTTG	TAGTCGGCTT	TGATTTCTAG	CTCCAGCTGG	8220

AATTTACAGCC	AATGAAGTAA	AGCAAGATGT	AACATCTGAA	GTGGTAATAG	GTGTGCTAGA	8280	
TTCTAAGGAG	GAATTGAAAG	AGTCAGAAA	TGATGCTCCA	AAACTAGAAA	CTCCTCTTAG	8340	
AGAGGAGCCA	AGACTAGCTC	CTCAAACGCT	TCCGGAAGCA	AGTGAAGTTC	TTGAAAACAA	8400	
AAGGGAAGAG	TCAAAAAGTAG	AGATAACATA	ACCAGCTCAA	GCGGATGATA	TCCGCAAGGT	8460	
TGTTGGGGAA	TTAGCCAAGG	ATATAAGTAT	TACTAAGTTG	TATATGACAG	GTCATTCTCT	8520	
TGGATGTTAC	CTAGCTCAGA	TTGCAGCGGT	TGAAGCTTAC	CAAAAATATC	CTGATTTTTA	8580	
TAACCATGTA	TTGAGGAAAG	TGACAACTTT	CAGTGCCTCCT	AAAGTGATTA	CTTCCAGAAC	8640	
TGTTTGGAAT	GCTAAGAATG	GTTTCTGGGA	TGTTGGTTTG	GAAAGTCGTA	AATTAGCTGT	8700	
TAGTGAAAA	ATTAAGCATT	ATGTGGTTGA	TAATGACAAT	GTTGTGACTC	CCTTGATTCA	8760	
TAATAATCGT	GATATTGTTA	CATTTACAGG	TAATTCACGC	TTTAAACACC	GTTCTCGTGG	8820	
CTATTTTGAA	AGTCCAATGA	ATGATAATCC	TAACTTAAT	ATTGGTAAAC	AAGCTACCTT	8880	
GGATAAACAT	GGTTATCGTG	ATCCGAAATT	GGATAAAGTG	CGATTCTTTA	AGAAACAGGC	8940	
TCTGCCTCGA	TCTTCTAGTC	AACCAAGCGC	TGAACCAATG	GAAAATATG	CCTCAGGAAA	9000	
ACAGGTTACT	CAAAGTTCGA	CAGCTTTCGG	AGGAGATGCT	AGAAGAGCTG	TGGATGGCAA	9060	
AGTCGATGGT	AACTATGGTC	ACAATCTGT	CACTCATA	CACTCATA	AACTTCCAAT	CTAAGCCTTG	9120
GTGGCAAGTA	GATTTGGCTA	AAGAAGAAAC	CATTGCGCAA	ATCAATATTT	ACAACCGAAC	9180	
AGACTGACC	CAGGATAGAT	TGGCAAACCTT	TGATGTCATT	CTTTTAGACA	GTTCTGGTAA	9240	
AGAAATTGAG	TGAAAACGTA	TAACATCTCC	TAAAGATGTG	TCAGCACAAA	TTACGATTAA	9300	
CCATAAAAA	GCGCGTATG	TTCGGATTGA	GCTAGAAGGC	TATAATGCC	TCAGTCTTGC	9360	
AGAAGTTGAA	GTTTTCTGCT	TTATAGCTAC	GAATGCTGAA	ACGGCGACAC	AAGTTTCTAA	9420	
GCCAGTTCAA	CCAATCAGTC	AGACTCCTGT	GAAGGATAAA	ACATTGACAA	TTCAACACAG	9480	
TGGAGCTTAC	ATTGCCCGCT	ACTCCATAAC	TTGGGAAGAA	GTTCCAGTAG	ATAAAGATGG	9540	
AAACCAAGTT	GTTCTGTAGTC	ATTCTTGGGA	AGGAAGCGGT	CGCAACCAGA	CTGCAGGTTT	9600	
TGTCTCAAC	CTCCCAATCA	AAGAAAATAT	GAGAAATCTG	CGAGTTAAGA	TTGAGAAAA	9660	
GACGGCCTA	CTATGGAATA	GATGGCAAAC	AATCTATGAA	AACAGACCAA	TTTTAGCTCA	9720	
ACCCACCGT	AAAATTACCC	ATTGGGGTAC	GACATTGAAT	TCCAAGGTGA	GTGACGATGA	9780	
TGTCTGTAA	TCTGATGGTA	GAATGACAGT	TAGTTTGTCT	AGTTTATAAG	AAAGTACTAC	9840	
CTGAGCTTGA	ATAGGACTCA	GGTAGCTCTC	TATGAAAGAA	CAAAATTAAT	ACTCAATGAA	9900	
AATCAAAGAG	CAAAC TAAGA	AACTAGCCGC	AGGTTGCTCA	AAGCACTGCT	TTGAGGTTGT	9960	

AGATAAGACT GACGAAGTCA GTCACATATA TAATCCAAGG CGACGTTGAC GTGGTTTGAA 10020
 GAGATTTTCG AAGAGTATAA ACAGAAAGGT AGAGCGCGTG TTCTAATTTG AACACGAGTA 10080
 GAAAACTTTT CTAAAAACAA AAACGAAAGG ATGGGTAAAC TGTATTCGCT GAACTGAATA 10140
 CGGGCGACTC TCCTCTAAAT CAAAATTAAG AAAGGAATTG ACCCCACCCT AAAAGTAGTG 10200
 GGAAAAAGAT AGTTGATCTA GCGAGCATCG CTCACTGCGC CCAACTCCTA TTTTCCCTTC 10260
 GCTTTTTGAT GGGTTTGTA TCTTCTCAA TATAAAATAT AAAATAAAGA AAGGTAGAGC 10320
 GTGTGTTTTG ATTTGAACAC GAGCGGAAAA CTCGGAAAAT AGATAATCTG ACTGAAAAAT 10380
 CAGGATTTCT CGTCAGGTC CTAATTTTCA GTCGTTTTCT TCTCGCTCTT TGTATCATAA 10440
 ATTATGTCTA TCCATATTGC TGCTCAGCAG GGTGAAATTG CTGATAAAAT TCTTCTTCCT 10500
 GGGGATCCTC TTCGTGCTAA GTTATTGCG GAGAATTTCC TTGATGATGC TGTGTTTTT 10560
 AACGAAGTGC GTAACATGTT TGTTTACT GGTACTTACA AGGGTCACTG TGTATCTGTC 10620
 ATGGGAAGTGGATGGGAAT GCCATCTATT TCGATTTATG CGCGTGAGTT AATCGTAGAC 10680
 TACGGTGTGA AGAAATTGAT TCGTGTGGGA ACTGCAGGTT CTTTGAATGA AGAGGTTTCA 10740
 GTTCGTGAAT TAGTTTTGGC GCAGGCGGCT GCAACCAACT CAAACATCGT TCGTAATGAC 10800
 TGGCCACAGT ACGATTTTCC ACAAATTGCT AGCTTTGATT TGCTTGATAA AGCCTACCAT 10860
 ATCGCCAAAA AACTTGGTAT GACTACTCAC GTTGGGAACG TTTTGTCAATC TGATGTCTTT 10920
 TACTCAAATT ACTTTGAAAA GAATATCGAG CTTGGTAAAT GGGGAGTCAA GGCTGTGGAA 10980
 ATGGAAGCAG CAGCTCTTTA CTATCTTGCT GCCCAATACC ATGTTGATGC GCTAGCTATC 11040
 ATGACCATCT CTGATAGCTT GTTCAATCCA GACGAAGACA CAACTGCAGA AGAACGTCAA 11100
 AATACCTTCA CTGATATGAT GAAGGTTGGT TTGAAACCT TGATTGCAGA ATAATTATAG 11160
 CAAAAAAGGG GCTCTTTGTC AACTGTAGTG GTTGAAAAA AAGCTAAGCT TGAGAAAGGA 11220
 CAAATTTTCT CTTTCTTTT TTGATATFCA GGGCGATAAA AATCCGTTTT TTGAAGTTTT 11280
 CAAAGTTCCG AAAACCAAAG GCATTGCGCT TGATAAGTTT GATGAGATTA TTGGTCGCTT 11340
 CCAGTTTGGC ATTAGAATAG TGTAGTTGAA GGGCGTTGAC GATTTTCTCT TTGTTCTTTA 11400
 GAAAGGTTTT AAAGACAGTC TGAAAAAGAG GATGAACCTG CTTTCAAGATG TCCTCAATGA 11460
 GTCCGAAAAA TTTCTCAGGG TCTTTGTTCT GAAAGTGAAA AAGTAAGAGT TGATAGATCT 11520
 GATAGTGGTG TTCAAGTCT TCTGAATAGC TTAAAATCTT GTCAAGAATT TCTTTATTTG 11580
 TTAAGTGCAT GCGAAAAGTA GGGCGATAAA AACGTTTATC GCTsArTTTA CGACTATCCT 11640
 GTTGGATGAG TTTCCAGTAA CGCTTGATAG CCTTGTATTC ATGAGATTTT CGTTCAAAC 11700
 GATTCATAAT TTGAACACGA AAACGACTCA TGGCACGGCT GAGATGTTGG ATAATATGGA 11760

AACGATCTAG AACGATTTTA GCACACGGAA AAAGCTGTTT AGCCAAGTCA TAGTAAGGAC 11820
 TAAACATATC CATCGTAATG ATTTTCACTT GACAACGAAC GGCTCTATCG TAGCGAAGAA 11880
 AGTGATTTTCG GATGACAGCT TGTGTTCTGC CTCAAGAAC AGTGATAATA TTAAGATTAT 11940
 CAAAATCTTG CGCAATGAAA CTCATCTTTC CCTTAGTGAA GGCATACTCA TCCCAAGACA 12000
 TAATCTTTGG AAGCCGAGAA AAATCATGCT CAAAGTGAAA GTCATTGAGC TTGCGAATGA 12060
 CAGTTGAAGT TGAAATGGCC AGCTGATGGG CAATATCAGT CATAGAAATT TTTTCAATTA 12120
 ACTTTTGAGC AATTTTTTGG TTGATGATAC GAGGGATTTG GTGATTTTTC TTTACCAGGG 12180
 GAGTCTCAGC AACCATCATT TTGGAAsAGT GATAGCACTT GAAACGGCGT TTTCTAAGGA 12240
 GAATTCCTAGA AGGCATACCA GTTGTTCGA GGTAAGGGAT CTAGACGGT TTTTGAAAGT 12300
 CATrTTTTCTT CATTAGACTT CCACAATCAG GGCAAGATGG AGCCTCATAA TCCAGCTTAG 12360
 CGATAATTC TTGTGGGTA TCCATATTGA TGATATCTAG AATCTTGATG TTTGGGTCTT 12420
 TAATATCGAG CAGTTTTGTG ATAAAATGTA ATGTTCAT ATGATTCTTT CTAATGAGTT 12480
 GTTTGTGCGC TTTTCATTAT AGGTCATATG GGACTTTTTT TCTACACAAA AATAGGCTCC 12540
 ATAATATCTA TAGTGGATTT ACCCACTACA AATATTATAG AGCCCAAAAA GGAAGCCCTT 12600
 TATGAATGT AGGACTTCTT TTTCTTATCC AGAAATTGAT CTAGCTCTCT CTGATTTGCA 12660
 AGAATAGTGA CTTTATGTGA ATATTCTTGG CAAAGTTTTT GGTAATTTTC TTTTGAGTT 12720
 TTGCGGACGC CCATCCCAAA GAATCCATCT GATAAACTCC CACTCAAAGC GTTCAGGGCA 12780
 ATCTACCGCC AFACTTTTCTC TGACTTTTCC ACGGTATTTA AGATAACGCT TAAAGGCTCT 12840
 AAAGAGACAG GTCAATGGCG AAAAATTGAG AAAGATGATT TGGTCAGCTT CTTGCATTG 12900
 TTCTTGGTAG TAGCACCAAG AATAATTACC ATCGATGACC CAAGCTTTAT GCTTGGTGAG 12960
 AAAGTTTTTT ATCTCGGTTA ACATCCATTC GCAGTCACTG TCTTGCCAAC CAGGTTGAAA 13020
 TTGGAGTGTG TCCATGTGCA GTTTTGGAAT GGAGTAGTAG TTAGATAACT TTTCTGCTAT 13080
 AGTTGACTTA CCAGAACCAG AATATCCGAT AATTGCGATT TTCATTTTCT ACCTTTTCCT 13140
 ATTTGGAGAC AAAAAACAG CCTCTATGGA CTGTTTCTTA TTTAACAAGT TTAGCTGAAA 13200
 GACGAGCTTT ATCGCGGCTT GCTTTGTTTT TGTGAATCAA ACCTTTAGTT TCTGCTTTAT 13260
 CGATAGCTGA GCTAGCAGCA CGGAAAAGTT CTTCAGATGG GTTTGCTTCG AAAGCTTTTA 13320
 TAGCAGTACG CATAGCTGAT TTTTGAGCTG AGTTCTTTTC GATTCGTCTA ACGTTCAATT 13380
 CAGCGCGTTT GATAGCTGAT TTAATGTTTG CCAATGGTCT TACCTCCATA TTTACTAACT 13440

(2) INFORMATION FOR SEQ ID NO: 129:

890

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 8512 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 129:

CCTTTTTC	AAA	ACTAGAT	ACT	AGTCTAT	CAA	AAGTAGG	AA	AGGGTTC	AAG	AAAATTG	60
ATTGGAAAT	TTTT	GAAAAT	CAT	AGAACTA	TT	AGCTAATC	CCT	AGTATTG	AAA	AGACTGG	120
ATAGCTTCT	TC	AGGTCATC	TT	GTAAC	TT	TCTCTGGT	CA	AGTTGGAC	AT	AGACTTCC	180
ACCAGACAGG	AT	CTAAAGTT	GG	AAAAATTTG	TAAA	AATCCT	CC	TTTCTTC	TAT	CGGAAAA	240
TCAACAGTTT	TT	ATCCAAGA	AG	CTACTTGT	TCT	TGCTCCA	ACT	TCCCTTG	TAAA	ATAGGT	300
TCATAGATCA	CT	CTTGCTAA	AC	GCCAATCC	TC	CATCATCTG	TAA	AGCGAAT	CG	CATCTCTT	360
TTAAATAGTT	GG	CCAAGTAT	AT	CAAATACT	TC	ATGAAC	TG	TTTTTAGG	AA	AGTCTGGA	420
TGACAAACCA	CCT	CTGTCAG	TAA	ATCGGCT	CC	ATGTGCAA	AAG	CGTGAAC	CCA	ACCATAC	480
TGACTTGAGA	AACC	CCTTGT	AT	CCTTTCT	TT	TGAAAGAT	AG	TGCAAGCC	TT	GATTTAAA	540
AGGACATTAC	GA	ATTCTCG	AGA	AGGATTT	CCC	AATGAT	CAA	ACAACCA	CT	GGATTTCT	600
TCCTGGTTAT	AAT	TTGGTTT	TT	CTCTGCT	AT	TTTTCTTA	GT	AAATCTTG	AT	ACATGGTC	660
AATACCTCTA	CAT	TTCTAGC	AA	CTGTCAA	AA	AGGCAGTC	TT	AAATGACT	CA	ATATTGAA	720
TTCTCAATTA	AAT	ACAATCT	GAT	ATAAAAT	GAC	GTAATA	ACT	ATCAATA	CC	AGTTCTAC	780
AGTAAGTTCA	AAT	TTAACAT	CAC	GACCTTC	AAC	GACATTT	TT	GAAAATAG	CT	ACAAC	840
GACAAATAGA	AT	GACGCTTA	ACA	AGCCCAT	AA	ACATCATT	CT	AAAAAAT	TT	TCTATTCC	900
CCTACTCTCC	CA	ACTCAGCA	CT	ATAGGAGA	TA	ATCTGGTC	AA	CTGTGCA	GAC	AAGAATT	960
GGATGGTATC	AC	GGAGTGGT	TT	GTCTGTTG	AA	ATATCAGC	ACC	GATAATC	AT	GGCTGACT	1020
CAAGTGGTGT	CT	TGCTACCA	CCT	GATTTGA	GG	AGATTGAG	CC	AGTCTTCA	GCT	CCAGTTT	1080
CAGAAATGTTT	TAG	ATGAAGG	TA	ACCAGCAG	TC	GAGATAAC	TAG	TCTGCT	GAG	TAAGTGT	1140
AACTATACAA	GCC	CATATAG	TAG	TGAGCTT	GG	CGATCCA	AG	TGAGAGTT	GC	ATCATCGT	1200
CAATTTCAAT	AG	CATCTCCC	CAG	AAATCCG	TCA	AACTTC	CT	TATAATG	CT	GTTGAGCT	1260
TGCTTGCTCC	AA	AGTCTCC	CCT	TCTTCAA	TCA	ATGTATA	CAC	CTTACGC	TG	GAAGCGG	1320
CTTCCAAGAG	GT	GGGTGATA	AAG	TATGGA	AG	TAGGTGTC	TG	TCAAGCGA	TG	AGCCAGAG	1380
CGAAGCGTTT	TT	GACGTGGG	TC	ATTAGACT	GG	TCTCCAA	GT	AATCACTG	AG	TAGCAATT	1440
CATGAAGGT	TG	ACGGTGC	TCA	ACATAGT	AG	GTCGACAT	AT	GGGCATTG	AAG	TAAC	1500

GATGATTGTC	TGAAAAGATG	AATTGACCAG	AATGCCCGAT	TTCATGAATC	AAGGTATAGA	1560
CATCGCTCAA	ACGGCCTGTC	CAGCTCATGA	GTACATAAGG	GTGTACGCGA	TATGGGTCCG	1620
CCGCATAACC	ACCGGAATCC	TTGCCACTGT	TAGCAGCAAA	GTCCACCCAG	CGCTCTTCTT	1680
GGTAACGAGC	AACTTCCTGA	CAATATCTTT	GCCCCAAAGG	TTCTACCGAC	TTCATGACCA	1740
AATCATAGGC	ATCGTCAATA	GTCACCTCAG	GATTCAGGGC	GCTGTCCAAG	TCCAATTTCC	1800
AGTCTGCAAA	GGTCATCTTT	TCAAGACCAT	TTACCTTGGC	AACATGCTTG	AGGTATCTCT	1860
GAGCGACTGG	TGCAAAGTCC	TTCATGATGA	GGTCAATCTG	GCGGTCAAAC	ATGACACGGT	1920
CCACTTCTTG	TTCAGCTAGA	AGATAGTCAA	AGACAGAGTC	GTATCCCTTC	ATATCAGCCA	1980
AGAGTTTTTC	AGACTTGACC	TGAGCCAGAT	AGGCTGCTGC	AGCCGTATTT	TGGTGCCTTAC	2040
GAAGTCCCTC	TGAGAAGGAA	CGGAAGGATT	TCTCACGAAC	CTCAGCATCC	TCATGGTTTT	2100
GGTAGAAAT	TTCATAGGTC	ACAAAGCTGT	TTTTGTAGGT	CTTGCCATGG	GCTTCAAAGT	2160
CAGCCATTTT	AAAATCCCA	GCTCGCATCT	TAGTATAAAT	GTCCTGCGGA	CTGTAGAAAA	2220
CTTCACCGAG	ATTTGTCAAG	GCCTTCTCCA	CATCTGCCCC	TAAGTAGTGG	GCTTTTTTGA	2280
TTTTAGCCTG	ACGAATGGCA	GCTGTTAAAT	GTGGCAATTT	ACCCAAACGG	TCCAAGACTT	2340
CCTCATCTGC	TGCCACCAAG	GCATCGTCAA	AGAAGGTCAA	GGCTACGCTG	GCATCTGTTT	2400
CAAATCCCAT	CCCAGCTTGG	GCAATATTGG	CAAATTCGTC	ATTGCTATAG	TCCGTCGTCT	2460
GAGGCATAAA	ACCATAGTTG	CCAATATGGC	TCATCTGAAT	GTAGATCTGT	TCCAATTCCG	2520
CAAAGCCTT	CTCGAAATCC	TCAAAAGTGT	GAAGATTGCC	CTTGTAATCA	CGGCTAAACT	2580
GGTTGATGTC	TTCGCGAGCT	TTCTCGATTG	CACGCAAGAA	ATCCTCACGG	TCTTGGTATA	2640
GGGCTGTAA	GTCCCAGAGT	TCCTTCTCTG	GAAATTCTGA	ACGGTGTTTT	TGTTCCATTT	2700
TCTTCCTCTT	ATTTCTCTAA	TTCTACTAAA	ACACTAAGGG	CTGATAAAGC	GTAAAGCGGT	2760
GCTGTTTCTG	CTCGCAAAT	ACGAGGACCT	AGGCCGTGCC	AAACGGCTCC	TTTAGCTTCA	2820
AAACTTTCGA	TTTCTGCAGG	TGAGAGACCG	CCTTCTGGAC	CAAAGATAAA	GAGCAGTTTG	2880
GCTCCTGTTT	CAAGACCAGT	GACTGCTTGC	AGAAGCGCAG	CGGCTTCTCC	TTCTTTAGCT	2940
GATTCTTCAT	AGGCTACTAT	GATAGAGTCA	AACTGGTCCA	GCTGAGCTAG	AAAATCTGCT	3000
TTTTTCTCGA	AAAGTTAAT	ACTTGGTACA	ATATTACGCT	TGCTTTGCTC	GGCTGCTCCA	3060
AGGGCAATTT	TTTCTAGTTT	TTCAACTTTT	TTACCCAATT	TCTTGCCATC	CCACTTGGCA	3120
ACTGACCAGT	CTGCAGGAAA	GGCCCAGATT	TGGCTAGCCC	CCAGTTCGGT	TACTTTTTGA	3180
GCGATGAACT	CCAGCTTGTC	TCCCTTGGGA	AATCCAGATG	CGATGGTCAC	TTGGACTGGT	3240

AGTTCCACAT	TGTCATTTAA	TTCTTGGACC	AACTCAAAC	GACGATTTTC	CATATCCAGC	3300
ACGCGCGCCA	AGCGCTTGAT	GCCATCATCA	AAGACTAAGG	TAACCTCATC	CTCTTCTTTC	3360
AAGCGCATAA	CCTGAAACAT	ATGCTTACTG	GTTTCCTTGT	CCTCGATAGT	GACAGGAGAG	3420
ATAGCACTGC	CTTTTACAAA	ATACTGCTGC	ATGCTAGCCT	CCAATCACAC	CAGAGATATC	3480
CTTGGTTTTC	TTAAAGACAC	AGGTATFCCA	TTCCCCTTGA	ACCATGTGAG	TTTCGAGGAA	3540
AAATCCAGCT	GAFTCAGCCG	ACTGGCGCAC	CATGTCCAAC	TTGTCCCTGA	TAATGCCACT	3600
CATGATCAGG	TAGCCTTCAT	CCTTTACCAA	GCGATAAGCA	TCGTCTATTA	GATGAATGAG	3660
GATATCCGCC	AAGATATTAG	CCACAATCAC	ATCTGCCTCA	ATTTCCACAC	CCTTAAGCAA	3720
ATCTCCAGCC	GCTACATGGA	TATTTTCCAT	GCCAGGTTG	AGCTCAATAT	TTTCTTGAGC	3780
CACACGAACC	GCCACATCAT	CCAGGTCATA	GGCGAAAATT	TCTTTAGCCC	CCAGAAGCGA	3840
GCTGGCAATA	GAGAGAACCC	CTGAACCAGT	CCCCACATCT	AGCACCCTTT	CGCCACCACG	3900
AAGAACCTGT	TCCAAGGCAA	AAAGGCTCAT	CTTGGTAGTT	GGGTGGGTTT	CAGTACCAAA	3960
AGCCATGCCA	GGATCCAGCT	TGATAATCAT	TTCCCCGCA	GTCGCCTCAT	AGTCTGTCCA	4020
AGAGGAACG	ATGGTCAAAT	CATGAGTGAT	ACGAGCAGGT	TCATAGTATT	TCTTCCAGTT	4080
GTCTGCCAG	TCTTCCTCAG	CCAAGGCAGT	CGTACCTATT	TTTAACTCTC	CCAAATCCAT	4140
AAAATCTGTC	AATTTCTGTA	GACGAGCCTG	CAAATCCGCC	TCAACCACTG	TCACATCCAC	4200
CGTGTGAGG	TAGTAGGCTG	TCACTACGAT	TTCTTCTTGC	TGCTCCACCT	CTGGGAAAAT	4260
CTCTCCAAAG	CGGTCCACAT	TTCCCACATA	GTCCATACTG	TCTTCGATTG	CGACTCCTTG	4320
CGCTCCCAGC	TCAATCAAGA	GATTGGAAAC	CAACTCCTCT	CCCTCACGCT	TCACTGTAAC	4380
TTTTAACTCT	TGCCATGTTT	CCATTATTTAA	TACCAAGCCC	GTAACACACA	AAACCAAAAT	4440
AGGAAATCT	CTGAAGACGC	TTGTGTCTAA	GAGAAGTTTA	TCTTTTTGGC	ACAGTGTTTA	4500
GGGCGGGTTC	AGTTTAGAAA	TGTAAGTCAA	CCATCCTTTC	TAATCACTTA	CTTTTAAATA	4560
ATCTTTAAT	CTCTCTGCA	ACTGAGGCAC	AACTTGACTG	GAACAAAGAA	ATTCCTCAAC	4620
ATTCATCAGC	TGATAGCCCT	GTCCTTCATC	TCCGAAGATG	ATATTGTCAA	ATTGTTCTTG	4680
TCTTAGCTGA	CCAACCATAA	AGACCGATTT	CTTGCCTTTA	AAAATTACGC	TAGGATAAAT	4740
CTTGCTCCAA	AGCAGACAGT	CTTCATCTAA	ATGAATTCCC	AGTTCCTCAT	AAACTTCACG	4800
CCGAGCGCAT	TCAAAAGGGC	TTTCGTCCCC	TTCACGGCCA	CCACCTGGCA	GTTCCACAT	4860
ATTGGCCAG	GGAATACTTG	CCTTATCATC	GCGTAAGATA	GTCAAAAAGCT	TATCCCACACA	4920
AAACAAAGCA	ATCTTGCAAC	CTGTGAAATC	AGAAATTTCT	AGTTCATCT	TCAGTTCCTT	4980
CTAACATTT	CTTTTCCAGC	TCGGCTAACC	AGTTTTCATA	ATATCTTTTC	TCATCCCTCA	5040

ACATTCGACT	ACTATCCATT	TTCTGTCTAG	CAATCTTGAG	AGCCTTACGA	GTTTCGATCTA	5100
CATCTTTCTT	CACCTTTAAT	TGATACCAGG	CTTGTATCAC	TTGAAGATTG	GACAGTTTGA	5160
GAGACAGAAA	CGATTTGACC	TGTCGAATAC	TAGCATATTG	CTCCGCTTGC	TCAAAAATCTC	5220
CTTCCAACAA	GGCGATATGA	AGCAGGGATA	GTTGGGCAAC	TGTCTGCATC	ATCGGAGTAG	5280
TTGTCCCTCTC	AAGTAATGCT	TGAAACTGCT	GTTTAGCTAC	TTCTTCCTTC	CCTTCCAAAA	5340
TGGAACTTC	ACCTTGCATA	CCTAATACAC	CATCCGCAAA	ACTCCCTCGT	GCATCCTCAG	5400
GAAGTCTTG	AACAAAGTCT	TTCAAATCAT	ATTCTTGAGG	AGCTAGCAAG	GTCTGGGCAG	5460
AATGTCTCAA	TACCAGGTAG	GCGTATTTGG	TATTTTCAGG	GTGTTGTAGT	AATTCCCAAA	5520
TTTTTGCTCC	ATCGGTGATG	TCGACTGGCA	AAATGTTATT	TAGGAAGAAA	GATAAATTAA	5580
GAAAAATCCA	AGTCCCTGCA	AAATACCAGC	TTCTTGTCAA	AAATCCAAAC	AATATCGCCA	5640
ATAATATCAA	GCCGAGATGA	ACCATCAAGC	CTCCTGAAAG	CATCAGGATG	ATTCTTTGAT	5700
CGCTTTCATC	CTCTTTTAAA	CCAATGTATT	GAGCACCAAC	ATTTTTCAGA	ATGGCTGTTC	5760
TACTAAGATG	AAACCTGCCT	GACTTTTGG	TCAAAATAAA	ATGTCCTAAT	CCAAAAGCCA	5820
CCAGCCGATA	GCCTGTCAAG	TAGCCACAAA	AAGCATGACC	CAGCTCATGA	AGAATAAAGA	5880
TTAAATACAT	GCTTAGAAGA	GCGAAGGCAT	AACCAAAAGT	AAAGGCTAAA	ACTGCGGAAT	5940
ACCCCAACTC	TGCAAATGCG	ATTGTTCCAC	AAGCAAAAGC	TAGCATAATA	AAGACAACAG	6000
CTAGCACATA	AACCAAATAA	GTCCCAATTT	TCTTCATAAC	ACCTCCAACC	AACTCCTAGT	6060
ATCTTGATA	AGGATAAAAT	TCTCCCTTTT	CCAAGCCAAT	TTTTCCCTCT	TCAAAGACTT	6120
CTTGGTTCCA	TTCCATGACA	AATTCCTCTG	CTTCTGGGTC	TTCCAAAAG	TCCATGAGGA	6180
CATCTAGCCC	AACCTCAGCA	GTATCTTTAA	GGAAAAGCGC	AAAATAAGCT	AAAAATTCAC	6240
GGGAAAATCC	TTTTTTAGGC	AGGTAAGGAA	TAACAGTCAA	ATAGTCTTCC	TCATTGACTG	6300
TTGACTTGGC	AGGATTGTAG	AAAAGGACCG	CTTCTCAAA	AAGAATGTCA	TCTGATGAAA	6360
CCTCTCCGTC	TTCATCCACC	ATCTCCACAC	CGCAGCATTT	TGCGCTTCCA	ATAGAAAAC	6420
CACTTCTACC	GCATGGTTGC	GTTTGTCCCA	GCTAATCTCA	AAGTCAAAGG	GAAAGTTCTT	6480
GTCCAACCTC	TCCTCTAAAA	TATCTAAAAA	TCCGTATGTT	GCCATTTTGT	CCTCTTTCTA	6540
TGCGACTCTT	TAATCGCCCC	GATTGCTCGG	AAATATGCTA	AAATAGATAC	TACCATCTTA	6600
CCACAAAATT	ATTTTATGTC	CTAATTATAC	CATATTACCT	CATTTAAACC	CTTGGTATCA	6660
GTGATTTTCT	TAAAAGTCTG	ATTTCTTCAT	TTCTCATAAA	AATCAATATA	AAAAGCCCTC	6720
GAAAGGGCTA	ATAAATCTAT	AAAATCAATA	GGCGAGTAAC	TAGCACAAGT	GGACGTGCTT	6780

TTTTATTGAC TATTACCACG ATACCACGCT TAATCTTAGG CTTGAACTTT CTTATCTGCA 6840
 ATAGCGTCTG TCAAAGTCTG AGAAAAGTTA AGCCCCATTT CTCGTCCCAA CTTATCTGCC 6900
 CATTTTGGTA TGGTCAAAGT CTTTTTAATG GGTTCCTGAC TTCCTAGGTA TTCTGATACA 6960
 TCAACAGATA CCATAGAAAT AAAAGATTTA TCAAGGTCAT AGGTTGACAC GAAATCTTCA 7020
 TCATCTTTAA AAGGATCATT ATCAATTAAA GACAAGCTAT TGATATCTGA TGGCTGAGGT 7080
 AACTCTCCAT CACTCTCTAT CAAATCTGCA ACAGTTATCC CTAGCCACTC CGACCCCAT 7140
 GCCAAAGCCT CAGAAATCCC CTCTCCTTGT GTAGCTGAGT ATTCAAAATC TGGGAAATGG 7200
 ACAAATAAG TCGCTTCTGT TCCGCTCTGTG TCGTCATAAT AAAATAAAGC TGGATACGTA 7260
 ACTAACATTT CACTACCTCC ATATCAAAAA GCAGGGACTG AATTTTACAA CCCAGCTTGC 7320
 TTTCTTATCC CTCTTTCAGT GTACTTATTC AGCTCACCAT GAAGGATTGT GATAGGTCTT 7380
 TCCCCTTGCT TTTCCATTTT AATATGGGAG CCTTTACCGC CTCTAGTCTT TATCCAACCA 7440
 TGGGCCGTAA GGAGTTTAA C ATCTCTTTT TGTGTCATAG GCATAGCGCT TTTACCTCCT 7500
 GACAACACCA TTATAACACG TGTTACACGT ATTGTAAAGG AGTGATACTT ATTATCTAT 7560
 TATACATAAA AGCCCCTAGA TGTGGTCTA AGGGAAGCCA ATTTATTCAT ACCTATTTTT 7620
 CTAATGAGTA GTAAAAACTG CTTCTTTATC GAGCAATTCA TCATCTGTAT AGTCAATTGT 7680
 AAAAGTATCT CGATCTAAGA CAGATTGAGG CGGAGTTGAA TGAATCATAG GAACACTGCG 7740
 TACTCTATAT TTTTATCTC CAATTTTAC AAACGATAC TCTTCGAAAA TCAAATTC 7800
 ACCACGTCAA CGTCGCCTTA CCGTACTCAA GTACAGcCTG CGGCTAGTTT CCTAGTTTGC 7860
 TCTTTGATTT TCATFGAGTA TGATTAATC TCAAGTCTC GAAATCAGGA TTTTCAACAG 7920
 TTATTACAAG GAGGCGATTT ACTACTTCAA AAACATCAAT TATTCATTTT TTCATATTTT 7980
 TTCAACCCAT TATTAGAATG AACTTCTTGG TAAGCAAAAT CAAGTTTAGA TTTAATGTTT 8040
 TCGTACAAAT CTAAAATCTC TTTTGGAGTA TCTTCCCGGA AGAAAAGTTT TCTTTTCCCT 8100
 GAAATAACTT GATCACTAAG AATCCAATGA CGAATTTGTT TTGTAAAAAT CAAAATTTCC 8160
 TGACTTGGTA GTTCCATCAT TTCCATTGCT TATCACCTCT CTTTTCATTA TAGTTCATAC 8220
 AATGACATTC AGCAATATTA TTTCTCAAGT CAGCACTTCC ACTTCTTTAG GCTCAACTAT 8280
 CCTATTTTGA GCTTTAAGGA AAATCAAATC TCTCATGCTG ATACCTCTCC TCATTA AATT 8340
 AAATAGTAAA AAAGATTCTA TCTCACTCCC TGATTATTAC AAAACCATTG AAATATCACA 8400
 ACTAATAGGC TAGAATGGAC ATAGTAAGAT ATAGTAGATG AGTCATTCTA CTC AAATCCA 8460
 CGTTAGAAAAG GACTGCTATG CCAGACAATC TCGCCGTTCC CATGCGCCCh GG 8512

(2) INFORMATION FOR SEQ ID NO: 130:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 2869 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 130:

```

CTCGTTTCAA GGTGAGTCT CTTGCAAATC TTGTTCGCGT TCTTCCTTTT GCCAAGGCAT      60
CTCTCCCATG GTTGGTGCCa GCCATTGTTG GAATCTTGCT CTCATTGGTT CTACCAAACA      120
AGCAAGAAAG CGATGTTTTT GAAATGGAAT AATCACTTAA ATCACTTTTG TAGCCAAGTC      180
TACAGGAGTG ATtkTCTTTT TTTATCCGAT GATAAATGTG TTATAATAGG TAGCGAAAGA      240
GGTGAAGAAA TGAATCAAAC AGTAGAATAT ATCAAAGAAC TGACAGCCAT TGCGtCGCCA      300
ACAGGCTTFA CTCGTGAGAT TGCGGACTAT TTAGTCAAGA CTCTAGAAGG TTTTGTTTAC      360
CAGCCGGTTC GCACATCCAA GGGCGGTGTC AATGTAACTA TTAAAGGTCA AAATGATGAG      420
CAACATCGCT ATGTGACTGC CCATGTAGAT ACGCTTGGTG CTATTGTCCG TGCTGTCAA      480
CCAGACGGCC GTCTCAAAT GGACCGTATC GGTGGCTTTC CTTGGAACAT GATTGAAGGA      540
GAAAACGTGA CCATTCATGT GGCTAGCACA GGTGAAAAAG TATCAGGAAC CATCCTCATC      600
CACCAAACCTT CTTGCCATGT CTATAAGGAT GCAGGAACTG CAGAACGCAC GCAAGACAAT      660
ATGGAAGTGC GTTTGGACGC CAAAGTAACT AGTGAAAAAG AACTCGTGC TCTTGGCATT      720
GAGGTCGGTG ATTTTATCAG TTTTGACCCA CGAACTGTCG TGACAGAGAC AGGTTTTATC      780
AAGTCTCGCC ATTTGGATGA CAAGGTCAGT GCGGCGATTT TGCTCAATCT CCTTCGCATT      840
TATAAGGAAG AGAAGATGA ATTGCCCGTA ACAACTCATT TTGCTTTTTC AGTCTTTGAA      900
GAAGTGGGAC ACGGTGCAAA CTCTAACATT CCTGCTCAGG TAGTAGAATA TCTGGCTGTG      960
GATATGGGAG CCATGGGAGA TGACCAGCAA ACAGACGAAT ATACAGTGTC TATCTGTGTC     1020
AAGGATGCTT CTGGACCTTA TCACTATGAC TTCCGTCAAC ATTTGGTGGC TTTGGCGAAA     1080
GAGCAAGATA TTCCATTTAA GCTGGATATC TATCCATTTT ATGGTTCGGA CGCTTCAGCG     1140
GCTATGTCTG CAGGGGCAGA AGTCAAACAC GCCCTTCTCG GTGCTGGTAT AGAGTCTAGC     1200
CATTCCTATG AGCGTACCCA TATTGACTCG GTGATCGCAA CAGAACGAAT GGTCGATGCT     1260
TATCTTAAGA GCACGTGGT GGACTAATAT GTGCCTTATT TGTcAGAGAA TTGACCTCAT     1320
CAAGAAGGAA GAAAATCCTT ACTTTGTCAA AGAGTTGGAA ACAGGCTATC TTGTGGTTGG     1380
AGACCACCAG TATTTTGAAG GCTATAGTCT CTTTCTAGCC AAGGAGCATG TCAGCGAATT     1440
    
```

896

GCACCATTTG AAAAAGGAGA CAAGACTCCG TTTTCTAGAA GAAATGAGTT TAGTCCAAGA 1500
GGCAGTTGCC AAGGCCTTGG CTGCTGAGAA AATGAATATC GAACTGCTAG GAAATGGCGA 1560
TGCTCATCTT CATTGGCATC TGTTCACG ACGGACAGGT GATATGAATG GTCATGGTCT 1620
CAAGGGTCGT GGACCAGTCT GGTGGGTTCC CTTTGAAGAA ATGACAGCAG AAACCTGCCA 1680
AGCAAACCG GATGAGATTA AAAGATTAGT CAAACGTTTA TCGTCAGAAG TAGATAAACT 1740
ATTAGAAATA AAGGAGTAGA AATGAAGAAA AGATACCTAG TCTTGACAGC TTTGCTAGCC 1800
TTGAGTCTAG CAGCTGTTC ACAAGAAAA ACAAAAAATG AAGATGGAGA AACTAAGACA 1860
GAACAGACAG CCAAAGCTGA TGGAACAGTC GGTAGTAAGT CTCAAGGAGC TGCCAGAAG 1920
AAAGCAGAAG TGGTCAATAA AGGTGATTAC TACAGCATTC AAGGAAATA CGATGAAATC 1980
ATCGTAGCCA ACAAACACTA TCCATTGTCT AAAGACTATA ATCCAGGGGA AAATCCAACA 2040
GCCAAGGCAG AGTTGGTCAA ACTCATCAA GCGATGCAAG AGGCAGGTTT CCCTATTAGT 2100
GATCATTACA GTGGTTTTAG AAGTTATGAA ACTCAGACCA AGCTCTATCA AGATTATGTC 2160
AACCAAGATG GAAAGGCAGC AGCTGACCGT TACTCTGCCC GTCCTGGCTA TAGCGAACAC 2220
CAGACAGGCT TGGCCTTTGA TGTGATTGGG ACTGATGGTG ATTTGGTGAC AGAAGAAAA 2280
GCAGCCCAAT GGCTCTTGA TCATGCAGCT GATATGGCT TTGTGTCCG TTATCTCAA 2340
GGCAAGGAAA AGGAAACAGG CTATATGGCT GAAGAATGGC ACCTGCGTTA TGTAGGAAAA 2400
GAAGCTAAAG AAATTGCTGC AAGTGGTCTC AGTTTGAAG AATACTATGG CTTTGAAGGC 2460
GGAGACTACG TCGATTAATA CTCTTCGAAA ATCTCTTCAA ACCACGTCAG CGTCGCCTTA 2520
CCTACTGACT GCGTCGGTTC TATTCACAAC CTCAAAACAG TGTTTTGAGT CGATTTCGTCA 2580
GTTTTATCTG CAACCTCAA GCTGTACTTT GAGCAstGCG GCTAGCTTCC TAGTTTGCTC 2640
TTTGATTTTC ATTGAGTACA AAAAGTAAAC TTTTCTCTTG CAATTCCAGA TAAATAGTGT 2700
ATAATGGATG GGTATGTGAA AACATACTT GTGGGAGGTA AAAATCTCTA ATTACCGCCA 2760
AAACCACAAA GGAGGATTTA AAAATGGCTA AAAAAGTCGA AAAACTTGTA AAATTGCAAA 2820
TCCCTGCTGG TAAAGCTACA CCAGCTCCAC CGGTTGGACC TGCTCTTG 2869

(2) INFORMATION FOR SEQ ID NO: 131:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 6186 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 131:

CTGAATCCCT TATAGGAGTC CAGTAACTTT TTAGCCTCTA CTTTGCCTTC ATAGGCAGCT 60
 TCAACATCAT TAAAAAAGA AIGCACTGAA GCAAGTTCCT CAGTGCTCCA CGACAAATCT 120
 AGTGGGTAAAC TATACTGTTT GTTCATTAAC TAATACCAGC TCTCATTCTT GCTTCTTTTA 180
 GTTCTTGCTT ACGATAACTA CGAGGGAGAA AAGCACGAAT CTCATCTTCA TTAAAACCGA 240
 TTTGCATACG CTTGCCATCA ATAATAATG GACGACGCAA AAGACTAGGA TACTGCTCAA 300
 TCAAATGAAG CAATTCCGAT ACCGAAATAC TCTCTACATC AATATTCAAT TTTTGAAAA 360
 TTTTTGAACG AGTTGAAATG ATGTCATCAG TACCATTTTC GGTCAAGGAA AGGATGTGTT 420
 GCAATCTTT TCTTGTTAAA GGACTGGTCA TAATATTGTG TTCCACAAAAG GGAACCTTATG 480
 TTTTTCTAAC CAGGCCCTAG CCTTACGACA TGATGTACAG CTCGGTGATA GAAATAGTGT 540
 AATCATGCTT TTCTCTCTT ATCTATACTT TGCTACTTCT ATTATACAAA AAAATAAAGC 600
 GCTTGACTAG GGATTTTTAG AAAAAAGCC TATTTTTTCA AGAAAAATAG GCTTTTTGCG 660
 AACGATTGAC ACAATTGGAT TTGGTTAATT CACTCTTAAC GATGGTTTTA AACGATATAT 720
 ATTTTTATAT ATGTAAATTA AAAACATCTT TCCTTTCACT TCCTACGACT TTTCAGATAC 780
 AGATAGCCAA AGAAGTTTTC ATAGAGGGCA AAAAAGAGGA GGAAGGCATG AAGAAAGAAG 840
 GTCTCTGGCA AAATCATAAT AACAGGATCC TTGGCTGGAT CAAAAGCCA GGTATCATCT 900
 CCCACAAAAG GAATTTGATG GAAAAGAGTA AAGAATTGGT CAAAACCAAT CAAAACCTCC 960
 CCAAGTCCAA TCATCACAGG TAAGACTACT AGAGCCAGGA GACTTTTTTCG ATAAAGAGAC 1020
 AAAAAGTCTT TTTTCACAAT CCTATTGACA AAGACATAGA AACTTGGCAG TGTCAC TAGA 1080
 GCTACTAGCT GAACCAAATG AAAGAGATTC TTGACCACTG CGAAATGGTG CAGACCAGCT 1140
 GCTGACGAAC GAAAATCAGG CATCTGTAAG ACCTGACTAA AAGGATTGGT CAGATAATTC 1200
 ATCAAGATAT GAAAATGTGA TTGAATGGTT TCTGGTTTTA GATAGACTCG ATTCGTTAAG 1260
 TTTAGCCACT GAATCTCCAT AGGATAGAAA ATCCAAGCCA GATAAATGGT CAGAAGGATG 1320
 GAGAGGGAGA GGAGAAAGAG CATAGAGCCC CAAAAGATCA ATTTAGTTTTT CATCAAAATC 1380
 CCACTCCGCA AGGCTAGAAA CCACATGTGT CGGTGCGATT GGCAGGCCAG CTACTTCTTC 1440
 TGCCTTAGTA AAACCTGTCTG TCACCAAGAG CGTTGGAATG CCATTGTCAA TCCCAGCCCG 1500
 AATATCAGTC AAATAATTGT CCCCAACCAT GATTAACCTC TCACGTTCCA AACCTAAGTG 1560
 CTCAACCGCC TTGTCCATAA TGATGGCATT TGGTTTTCCG ATATAAACCG GCTTCACTCG 1620
 TGTCGCTACT TCAAGCAGCG TAATCAGTGA GCCAGCACCT GGCAAAAAGAC CGCGTTCCTG 1680
 CGGGATGTTG AGGTCAGGAT TGGTTCCGAT AAAATGGGCA CCCTTTTGAA TAGCAAGAGT 1740

TGCTGTGGCA	AATTTTTCAT	AGTCGACTTG	CCAATCCAGA	CCAACCTACCA	CGTAGGCAGG	1800
TTTTTCCTTG	TCTTCCACAT	AACCAGCCGC	CTTGATGGCT	TCCTTGAGTC	CTGCTTCTCC	1860
GACGACATAG	ACGGTCTTTT	CAAGCCCCAA	ATCATTCCATA	TAGTCGATGG	TTGCCAAAGT	1920
CGCTGTGTAG	ACAGTCGATA	GGGGCGTATC	GATATTAATA	TTCTGAGCCA	ACATCTCCTT	1980
AACACTCTCT	GGAGTGCGGG	TTGTATTGTT	GGTTACAAAG	AGATAGGGAA	TGTCCCGCTT	2040
TTGCAATTCA	TGAACAAAAG	TCTCTCCAGC	AGGGATTCCG	TCTTTCCCCT	TATAAATGGT	2100
TCCGCTAAA	TCAATTAAAT	AGCCTTTATA	TTTTCATCTAT	TTCTCCCTAA	GCCTTTTTTA	2160
TTTCTTGCCA	AGTAATGATT	GCTTGGGCAT	TGATAACCCC	ATCACTTGTA	ATTTTCATGCT	2220
TGCTTTCCAG	TCCAGTCCGT	TCAACAGCCG	ATGTAATCAC	CCCACCTGGT	CGAACTTCCCT	2280
TGACATACTT	GAGGTGATT	TTCTTGGGAA	TATAGTGGGT	CAAAAAATCC	GCTCCCATGA	2340
CCTCAAAAAT	CCAGTCCAAG	TATTTACTGT	TATTGACATG	ACCATTCCATA	TCCAAGTCGT	2400
AAAAACGAAC	ATGGTAATCC	TTGCTGATCG	GTTCTTCCAA	GGACTCATA	TTCGGTCCAC	2460
GGATAAGTTT	TTTATCAAAA	TCAGACTGGT	AAGGAGCCAC	AATCTCAGGT	TCAACAACAT	2520
GGACTTTTCG	ACTGTCGCGG	TCCATGAGAA	CAAAGGTCGC	CATCATGTGG	ATGAGCTCCT	2580
GCTCCGCTTC	ATTATAAATA	GTAAGCGAC	GGTAGCAAAA	AAGTCGATTG	TAGCTCAAGG	2640
CTTCCGTTTC	GATGGTAATT	TCTTCCGCAA	AACGAGGCAA	ACGAACCACC	TCAATATCAT	2700
ATTCTACGAT	AATCCAGACC	AGATTATATT	CTTCCAAAAT	GGCCTTATCA	CTAACTCCCA	2760
GTTCAATCGA	CTGCATCCCT	GAAACTTGCA	GTGACAGCAA	AATCACATCT	GGAAGTTTGA	2820
TATGACCGTT	CATATCAGCC	ATATCAAAAAG	GAATTTTCAT	TTTCATTTGA	TAAGTTAAGC	2880
CCATGATCCT	ACTCCAAAAT	AAATCGTTCT	GCTACAGTAT	CTCCCAAAA	GAGACCTCTC	2940
TTTGTTCATGC	GAACGTGGTC	ACCCTCAATC	TGCATGAGGC	CTTGTGTAAC	CAAATCTCTG	3000
ACAATTTCTC	CATAAAGTCC	AGCAAAAGAC	TGTCCAAAAT	TTTCCTCAAA	TCGCGCCATG	3060
GAAACCCCGG	ATTTCTTGCG	GAGTCCCAAG	AACATTTCTT	CTTCCATTTG	CTCCTTTTGA	3120
CTCAGGTGAT	CTTCTGTAAT	ACAAGCATTG	CCTTCCTCAA	CCGCACTGAG	ATAATGACGA	3180
ATGGGACCAT	GATTTTTATA	GCGTACTCCA	TTGACATAAC	CAGATGCCCC	TGCACCAATA	3240
CCATAGTATT	CAGCATTGTC	CCAGTACATG	AGATTATGAC	GACTTTCAAA	ACCGGGTTTG	3300
GAGAAATTAG	AAATCTCATA	ATGCTCAAAA	CCCGCTCGCT	CCAGCTCTGC	AATGATGTAC	3360
TCAAACATCT	CCGCTTCTAG	TTCTCCTTA	GGCAGAGGCA	ATTTCCACG	TCGCATCCGG	3420
TTCATAAAGA	CCGTATGGTT	TTCTAAAATC	AAACTATACA	AACTCATGTG	GGGAATATCC	3480
AATCCAATGG	CTTTAGCCAC	ATTTTCCTTT	ACTTGCTCCA	TGGTCTGACC	AGGCAGAGCA	3540

TAAATCAAAT	CAATGGAGAT	ATTGTCAAAA	CCAGCCAGTT	TCAGGCGATC	GATATTTTCA	3600
TAAATATCCT	TCTCCAAATG	ACTGCGCCCA	ATCTTTTTTCA	ACATCTTATC	ATCAAAGGTC	3660
TGGACACCTA	GCGAAACACG	ATTGACAGCC	GAATTTTTTCA	AAACAGCTAT	CTTATCCGCA	3720
TCCAAATCGC	CTGGATTGGC	TTCAATGGTC	AACTCTTCCA	AGACAGACAA	ATCCAAGTTT	3780
TTAGTCAAGC	CATTTCAGTAA	CACCTCCAGT	TGCGGAGCCG	ACAGGGCTGT	CGgTGTCCA	3840
CCACCGATAT	AAAGGGTTGA	CAACTTTTTCA	ATATCATAAG	AACGAAACTC	TTCCAGCAGA	3900
TGCTCTAAAT	AGCTGTCGAC	TGGCTGATTT	TTGATGAAGA	CCTTTGAAA	ATCACAATAA	3960
TAACAAATCT	GGGTACAAAA	TGGGATGTGC	ACATAGGCTG	ACGTTGGTTT	TTTCTGCATA	4020
GTAATTATTA	TACCACAAAG	ACTAGATTCC	AGATAAAAAT	CACCATCCCC	AGATACATAG	4080
TCCGTCCGGA	GATGGTGATG	GTTTATCTCT	CTGTTATATC	AATCACAATC	TCTTCTGAGT	4140
CATCAAGAGC	TTCGGCTTTT	TCTTGCCATT	GCTCCTGAG	ATTATTTAAT	TGATTTTTTG	4200
ATGCTTCTGT	CGCTTGAAAA	GCATAGGATT	TAGTTTGAGC	AAGTATACTG	TCCACAGTGA	4260
TTTCACCTGA	CTCAACCTGT	TCTTTTGTTT	TCAGAACAAA	ATCTGTAGCC	TGCTCCTTAA	4320
CTTCTGTCAG	TTTTTCACAG	ACTTGCTCCT	TGGCATACTC	CGGATCTTCT	CTCAAATCAT	4380
CTAGAAAATC	TTGAGCCTGA	CTGCAAACCT	GTTTGCCCTT	ATCACCTGTT	AAAAACAAGG	4440
CAAGAGCTGC	ACCTGAAACG	GTTCCATAAA	GGATTGAGGA	TAATTTACCC	ATAAGGATTC	4500
TCCTTTTTTA	TTTTTTGAAA	AATTTACTTG	CAAGACGAAG	AGCTGACAGA	CTTGCAACCAG	4560
TCTTGAGTGT	TTTTGAACCA	GCTGATGAAG	CTTCTTGCT	CAAGACACGC	GCATGGTCAT	4620
TGAGGTCTGA	AACAGATAGA	GATAAATCTG	CAACAGCACT	GAAGAGTGA	TCAATCGTAG	4680
CCACCTTGAC	ATTGATATCA	TCTGCCAAGA	CATTGACCTT	AGCCAACAAC	TCATTGGTGT	4740
GATGCAAGGT	CACATCCACA	TCTGAAGTCA	AGGTTTAAAT	CGTCTTTTCT	GTTTCATCGA	4800
TGACACGACC	AAGCTTTTGT	ACAGTAATGA	TCAGATAGAC	CAAAAAGACA	ATCAAAGCTA	4860
GGGCAACAAG	AATATATGCA	ACTTCTAACA	TTTAGTTTTT	CTCCTCTGTA	ATATAGTAAG	4920
GGGCCTTCTT	TCGATTTTGA	TAAATAACGA	TCATTATACC	GAGACCGATA	AGGACAACCTG	4980
ACAGCCATTG	GGACACTCGA	AAGCCGAAGA	ACATGAGACT	ATCTGTTCGC	ATACCTTCGA	5040
TAACCATACG	ACCGAAACCA	TACCAAATCA	AGTAAAAGGC	CGTGATATGA	CCTCGTCTGA	5100
GACTCTTCCA	TTTCCGTCTA	AAAATCAGAA	TCAAGGCAAA	GCCAAGCAGA	TTCCATAGAG	5160
ACTCATAAAG	GAAAGTCGGT	TGACGGTAGC	TCCCCTCAAT	ATACATCTGG	TCACGGATAA	5220
AGCCAGGTAG	ATAATCCAGA	TTATCCACTG	TTGCACCATA	AGCTTCTTGG	TTAAAGAAAT	5280

900

TACCCCAACG CCCCAAACCTT TGAGCAATCA TAACGCTAGG CGCCGCAATA TCTAGAAAAT	5340
CCCAAGTATT GATGAGTTTA CGGTACAGCAA AGATATAGAG CACAAGAGCC CCAGTTATCA	5400
AACCACCGTA AATGGCCAAA CCACCATTCC AAATGGCCAAA AATCTCTCCT AAATTCTGAC	5460
TATAGTAATC AAATCGGAAA ATAACATAGT AGAGACGAGC TCCTAAAATA GCCAAGGGAA	5520
AGGCTACTAA GATAAAATCT AAAATATCGT CTGGTATGAT CTTCTTTCTA GGTGCTTCTT	5580
TCATGGTCAA ATAAACCGCA AGAATCAAGC CTGTCAACAAT ACATAAGGCA TACCAACGAA	5640
TGGCTAGGGG TCCTAGTTGA ATAGCAATTG GATCAAGCAT TTTGCACCTC ATTTTCGAGCG	5700
ATTAGACTTG TCAGTCGTTT GTCGAACAAA CGGGTCGCAT CAAAGCCCAT TTCCTTGGA	5760
CGATAATFCA TGGCAGCTGC CTCAATCACA ACAGAGATAT TACGACCTGT TTTAACTGGA	5820
ATACGAATAC GAGGAATGtA CGCCAGAAAC TTCAAGTTCC TCTGCATTAT TTCCAAGACG	5880
ATCAAAGGTC TTATGCGTAT CGTAATTTTC CAAATAGACA GCAAGCTGAA CCTGTGAAGA	5940
ATCCTTGACA GCACTCGCAC CGTAGAGACT CATAACATCG ATAATACCAA CCCCACGAAT	6000
TTCAATCAAG TGTTTCAAAA TTTCAGCTGG TTCACCCAG AGAGTAATCT CATCCTTGGC	6060
AAAGATATCG ACACGGTCAT CGGCTACCAA ACGGTGACCA CGTTTGACAA GCTCAAGACC	6120
TGTCTCGCTC TTACCAATTC CACTATCTCC CTGAATCAAG ACGCCCATCC CATAAATATC	6180
CATCAA	6186

(2) INFORMATION FOR SEQ ID NO: 132:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 9541 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 132:

GAAAATCACA ACCCTTTTGG CAAAATTTTT GAGATTATTT TCACAAACTT GATTTTTCAA	60
AGTATACTCA ATAAAAATTA AAAAAATCCA CTACGTCAAG GCGAGGCTAA TGTGGTTTGA	120
AGAAATTTTC GAAGAGCGTG AATGAGTATC ATCTATAGTA AAATAAAAAA ACTGAACAAT	180
TTGGTTGGGG ACAGCCAAAC CAATTTCTCA CAATGTTTCA GAAACAAGGG TGTGCTATTC	240
CAATTCAGC CTAATAAAC TGTCATAGAT TGCTGAAACA AAGCTAGGT AAAAGTCTTC	300
ATAATAAAAA GACCTCCTAT CAAGTGTTC AAACTTTGA TAGGAGGTCT TGTTTTGTGA	360
AAATATTTAT CAAATTTTCT ATACAAGTGA GCTGTTAGCC AGGTTCTTTC TATTCTTTCA	420
ATTTCAATGA ATGGATTTTT TACTAATACT CATAACTGGG AATTTGCTCG TGTAATAATA	480

901

GCGAGATAGA	TGGTATTTAT	AAAACACTCA	AGACAGCTAG	ACTAATATCA	TTTAAAACAT	540
TATCTTCTTT	TGAGCGACTG	TTGGTTACCA	ACATAGCTAA	ATTTCCCTGCA	TTTTCAAATF	600
GATAGGGTTC	TGATTTAGCA	FTCACAACCA	CCAAGAGGTG	TTCTTTGCCG	TGAAC TTCAT	660
AGATAAGGTA	GCCGCTATGT	TCAATCGCAG	AATGCACAAA	GACATGATGG	TAAATTTTCAT	720
CATAGCTAGA	GTAAGAAAAG	GCACCAGTTT	TTGTCTTCAA	TCGGATGACT	TGACGGATAA	780
ACTCAATACT	GTCTTGACGC	TCATTAATCA	AGTTCCAGTT	CACTTGGTTC	ACACTGTCAG	840
GAGCATTATA	GCTATTCATC	GCACGCTCTC	TATCATCATG	GGTCAACTCA	CCATTTTCAC	900
CAGTCGCAAC	CAGTTTGGTA	CGACCAAATT	CTTGACCGAT	TCCATAAAG	GCCATCCCTT	960
GCATGAGCAG	ATTCATGGCT	GTGGCAGTTT	CGACCTTGCG	CATGATTTGC	TCTGAACTTT	1020
GGTCTGGATG	AAGGGTTGCC	AATAAATCGT	GAAGATTGTA	ATTGTCATGG	GCTTCTACAT	1080
AGTTAAGCAC	CTGATTTGGA	TGTGTATAGC	TTCCTAATTC	ACGACTTCCT	AGGATTGCTT	1140
TAGCTAGAAT	TGGCTCTGTC	GCAGCACCAC	TGACAAAACC	TGACTTGATA	GCACCATAAA	1200
CTTCTCCCCC	TTTGACAGCA	TCGCGCTGAT	TGTCATPAAA	GAAACCAATA	TTTGGCATCT	1260
GGTAGGCATT	GTCCTTCTTG	GCCTTATCAT	AAGGGGCAAG	ACCTGTTCCT	ATATCCCATC	1320
CTTCTCCATA	GAGGATAATG	TTGGAGTCGA	TTTCATCCAA	GCTTTGACGA	ATCATCTGCA	1380
TGGTCTTGAC	ATCATGAATC	CCCATCAAGT	CAAAACGGAA	GCCGTCAATA	TTATATTCTT	1440
GCACCCAGTA	TAGAAGAGAA	TCAATCATAT	ACTTGCGAAA	CATTTCTGTT	TCACTGGCTG	1500
TTTCATTTCC	AACACCCGTT	CCATTCTGGA	AGGTACCATC	TGGATTGATA	CGATAATAGT	1560
AATCAGGGAC	TGTTGTTTGG	AATGGTGCAT	CAACAAC TGA	GAAGGTATGG	TTATAGACTA	1620
CATCCATAAT	GACTCCAATA	CCCGCATCGT	GATAAGCTTG	AACCATCACC	TTCAAATCAC	1680
GAATGACCTG	AGCTGGATCA	TCTGGATTAG	TTGAAAAACT	AGTTTCTGGC	GCGTTATAGT	1740
TTTGTGGATC	ATAACCCAG	TTGTAGGTTA	CATTTCCATC	CTCATCGTAT	TCTTTATCAC	1800
GGTCTGCAAT	TGGTTGCAAT	TGAACATAAT	TGTAGCCCAG	CTTCTTGATG	TAATCAAAAAG	1860
CAGTTGACTG	GCCGTATTTG	TTAACTGTTC	CAGCCTGAGC	AGCACCCAAG	AAAGTTCCTC	1920
GAAGATGTTT	ATCTACACCC	GATGTAGGTG	ATTTAGTCAA	ATCACGAATG	TGCATTTTCAC	1980
AGATAACTGC	CTTACATGGA	TTTTCCAAGC	GCCAAGTAGC	CTCCGAACCG	TGCTTAACCT	2040
CGAAGTTTTC	AACTTGCTTT	TCTACATGGC	TCAGAATAGC	TGAACGTTTG	CCATCAGGGC	2100
TGGTCGCGAT	TGTATAAGGA	TCACGTGTCA	GTGTTTGGTG	ATGAGGGAAT	TGGACTTGAT	2160
ACTGATAAGT	CTTACCTACC	AAATCTTCTT	CAACATCCAA	ACTCCAGACA	CCGATTGTAT	2220

TGTCCTTATG	ATTATAAGAG	TAGCTATTGC	CTCTTTTCAT	CTCAAAAGTC	TTCCAAACGG	2280
GTGCATCATT	AGCAGCTGAT	TCATAAACGA	CAACTTGCAC	TTCTGTCGCT	GTAGGTGACC	2340
AGAGAGAAAA	ATGAGCCTGA	TTGTCTCTA	CACGGCAACC	CAATTCTCCT	TGGTAACCCC	2400
AATGATGATC	AAAAC TAGCA	CTGTTAATGG	CCTTATCAAA	GGCAAAAGGA	TTTTGATTTT	2460
TATAGAAAGG	ACTGGCAATA	GCAGGATTTT	CAGAGTAATA	AATCCTATCA	TCGCCTTCCA	2520
AAATCCAGAC	CTCTGTTAAT	AGGGGATAGT	GATTA AACG	GATAGAATAT	TCTTTACTAG	2580
TTTGACCTGT	ATGAACCACA	AAATTCAAGC	TTTCTATAAC	ATGTGAACTT	GGGTGTTCAA	2640
AGCTAAATAA	AGCTCCAAAA	TAATCTTCTT	TGTAGGTTAG	CAAATCAATT	CGTTGATCCT	2700
GACTTTTAC	AAAGGAGCAA	GTGTCAATAT	CTCCATCTT	ACGATGGTAA	TGAATGCGCA	2760
TAGGTTAGTT	ATACATTTTT	TATTTTTCCT	TTTTACTTTG	TTTCTATTTT	ACTAATAAAT	2820
TTTTGTCAAT	CTCGTCTCAA	TTAACAGACA	TAGTCATATT	CTCTAAACTC	TGTTTTTAAA	2880
CGATCCATTA	CAAAC TTTCT	AGCCATGCCT	CATCTCTGAC	CTGGATACCA	AGTTCTTGTG	2940
CTTTTGCAG	TTTACTTCCA	GCGTCTGCAC	CTACCACGAC	GAGGTCCGGT	TTTTTAGAAA	3000
TACTACCTGT	CACTTTGGCA	CCCAGACTTT	CGAGTTTACT	TTTAGCTTCT	GAGCGCTTGA	3060
GTCGTTCCAA	TTTTCTGTG	AATACCACGG	TCAAACCTGA	CAAGGCCGCA	TCCGCTACTA	3120
CCGTCTGTCC	TTTATAGTCC	AGATTGACCC	CAGTTTCTTT	CAATTC TCTG	AGCAGAATTT	3180
CAGAGCCTTC	TGTCGCAAAA	TAAGTCTGAA	GACTTTTGGC	AATCACGCCA	CCTAGACTTT	3240
CAACTACTAGC	CACTTCCTCT	GAATCTGCCT	GAGACAGATT	TTCAATTGAA	TGGAAATATT	3300
GAAGTAAAAG	CTGACTAACC	TTGCTTCCGA	CATGACGAAT	TCCCAAACCA	AATAAGAGCT	3360
TCTCGGCAGA	ATTTTCCTTT	GATGCTTGGA	TAGCCTGATA	CAGTTTAGCA	GCGGACTTTT	3420
CCTTAACTCC	CTCTAAAAGG	AGGAAATCCT	CTTCTTGCAA	ACGATAAATA	TCCGCCACAT	3480
CCTTGACTAA	ATTAGCAGCA	AAAAGCTTCT	CAACAATAGA	TGGACCAAGG	CCTGTAATAT	3540
TCATAGCATC	ACGAGAAGCA	AAGTGAATCA	AGCCTTCCAT	GATTTGAGCA	GGGCAACGCG	3600
GATTGATACA	ACGTAGGGCC	ACTTCATCTT	CAAAGTGCAA	CAAGTCAGAG	TTACAAC TTG	3660
GACAGTTTGT	AGGGATATCT	AGTTTTTCTT	CAGAAACCCG	TTTGGACTCT	ACCACACGTA	3720
AAACGGCAGG	GATGATGTCA	CCAGCCTTAT	ATACAATGAC	CGTATCGTCT	TTTCGGATAT	3780
CTTTTTCAGC	AATATAATCT	ACATTGTGCA	GGGTCGCACG	GCTAACAGTC	GTACCGGCAA	3840
GTTGTACTGG	TGTTAGATTA	GCAGTTGGAG	TTACAACACC	GGTACGGCCA	ACTGTCCAGT	3900
CAACTGATAA	GAGTTGAGCT	TCTTTTCTT	CGGCAGGGAA	CTTG TAGGCT	ACTGCCCACT	3960
TTGGAGCCTT	AACTGTAAAA	CCAAGTCTT	CTTGACTTGC	TAGGTCGTTG	ACCTTGATTA	4020

CCACTCCATC AATATCGTAA GGCAGATTTT CCCGTTCCCTG TCCTACTTCT TGGATAAAAT 4080
 TCCAGATTTT ATCTATGTTT TCAGCCAAGA TTCGCTTAGG ATTGACCACA AAACCTAGTT 4140
 GTTCTAGGTA CTTCAAACCC TTTTCTTGGC TATCACGAGT TGAAGGGCTG GCTTCTTGAT 4200
 AGAGAAACGT TGCAAGATTA CGCTTGGCAA CTACTGCTGT ATCCAAGTGA CGCAGAGTTC 4260
 CTGCTGCCGC ATTACGAGGA TTAGCAAATF CAGGCTCTCC ATTTTCTTGG CGCGCTTGGT 4320
 TAACTTGGTC AAAGGAAGCG CGTGGCATGT AACATTCCCC ACGAACTGTG ATATCTAGTT 4380
 CTTCTGGCAA AGTCAAAGGG ATGTCCTTAA CACGCTTGAG GTTTTCTGTG ATATTTTCAC 4440
 CAATTGAACC ATCTCCACGT GTTACCCAG CAACCAAAAT CCCCTTTTCA TAAGTCAGCG 4500
 AGATAGATAA GCCATCGATT TTCAGCTCAC AAATATAGGT CGGATGAGCC ACTTCCTTAC 4560
 GAACACGCGC ATCAAAGCA TCTAGCTCCT CACATGAAAA AGCATCCTGC AAATAATAA 4620
 GAGGATACTG ATGACTGTAT TTTTCAAAC CATCTAAAAC CTTGCCACCA ACACGATGAG 4680
 TCGGACTGTC TGCTAGCACT TGCTCTGGAT AAGCAGTTTC TAACTCGACC AACTCACGGT 4740
 AAAGGCGGTC ATACTCACTG TCTGAAACCG AGGGATTATC GCTGGTATAG TACTCAGTCG 4800
 CATAGCGATT GAGCAAAGCG ACTAACTCAT TCATTCTTTT ATTCATAAGA CCATTTTACC 4860
 ATAAAACAAG CCCTCCTCAC AAACGAGAAG GCGGAAAAA AACTTAGTT TGAATTATT 4920
 TTTGAAATC AAGCAACCTT ATATCAATTT TTCAAATGA GTTCGAACAT ATCCGAGAGC 4980
 TAAGAAATAT AAGGCTACAA CTCCAAGTCC AATAATCAAG AAAGAATAAA GATGGACACT 5040
 TGGCAAGACT GTCATAAATC CTTTGTCAAT AGGCATAAAT AGAATAGCTA AGGTAAAAAT 5100
 TGTAATATTA AAAATCGTCA TAAACAATTC ACAAACTAAA TTTCCAGAAA AGGAAAGAAA 5220
 AGTTGGAAGT GGTAAATCCCA TCATAAAAAAC TCCGACACCT GTCAAAGCCA GTAAAAATCAA 5280
 AAGATTATAA ATATTAGCTT TAATTTTACT AGCTAGAAGA GCCCCAATGA TGGAACCAAT 5340
 AGCCCCATA GTTAAAATAC TTGCATAGGC TCCTTCTGAC CCGTAAAGCT GATTCGAAAA 5400
 GGAAGTAGA AATTCAAAG CTGCAAAAAA GAAATTAACG CTGGAAGCTA CCAGCAAAAG 5460
 GAAGAAAATF TCTTGCTGAT GCCAGATATA GTGTAACCCA TCCTTGATAT CTACAAAAAT 5520
 ATCTCTCCA GTAAAAGCCT TTTTCTCTTG AACTTTTGCT TCCTCTTTTG GAAGGAAAGC 5580
 CACTAGAACA AAAGCAATGA AAAAAGTCAG CGAGTCTAGC AGTAGCGTCA TATGGAGACT 5640
 TGCAAACGTG AAAACAAGGA AGGAAAGAAC AGGAGAGCTA ACACCTACAA CCTGCAAAAC 5700
 CAGCTCTAAG CGAGAATTAT AGATCACAAAT CTCATCTTTC TCCACCCTT CAGTTATGAT 5760

AGCTTTATTG	GCTGTGCGAG	AAAAGGCAAA	AGCAATAGCC	TGCACAATGT	TAGCAACAAT	5820
CAAAGCGCCA	ATCATCCAGC	TATCATTCCT	TATGAAAGAA	ATAGCCAGAC	AAAGAATCCC	5880
ACAAACAAGA	TCTGCCGTCA	TTAAAATCTT	ACGACGAGAA	AAACGGTCTG	AAATAACTCC	5940
GCCAAAGGGA	TTGACGAGAA	TAGATGTGAC	GAGCTCAGAA	ATCTGATACA	TTCTTAAAC	6000
TGTCTGTCTT	ATAGTCCCCA	TAGAAGCCAA	CCAGACACTA	TTTCCATAAT	CATAGAGCAT	6060
ATTTCCCAT	TTATTGATAG	CCCCACGGCT	AATCAACTGC	ACTGCATAGC	GATTCATATT	6120
AAAGCTCCTC	TCAAATTTTG	AAACTATTGT	ATCAAAACCG	AAAGGAGCTT	TTTATTTT	6180
CCCTTATTTG	GGAAAATTAA	CTTTTGACAA	ATTTTTCGTA	GTGTTCCCTGA	TAATAGGCTA	6240
CTTGCTCTGG	AAGACCTAAC	ACATCAAAA	TATGCATGGC	CTCTTGCAATC	TGCTTACAGC	6300
CTTCTTTACA	CTGTCCFTTT	TGATATAAGG	CAAAACCTTT	TAAATAATGG	AAAACATTAC	6360
GCTCATAAAG	CTTAATACCT	TTGTCAATAA	TCTTCTCTGT	ATAAGCCTCA	AAATAGTTGG	6420
CATTATAAAA	AGAAGAATGC	TCTAAACAAT	GCTGGTAAACA	ATTGAGGGCC	AAAATCAACA	6480
CTAATCTCTT	ATGGCGACTA	ATCTCTTGGT	AAAATTCCTC	CCTCTCCATA	ACTTCTCTAC	6540
CAATCCGAGT	GACATAGTCT	ACATCGTAGA	AACTATAGAG	GTTACCGAAA	AGAATCAACT	6600
CATACATGGT	CCATTCCTTCT	GTTTTGAAGA	GATAATCTGC	TACCTTACCC	AAATCATCCT	6660
GCTTCATATC	ATAACTCGCA	TCTCTTTGAC	AAATCAGACC	TTGTAGCAAA	ATCCAGTTCA	6720
GCTCAAAATA	AAGGGGAGTC	GTCGAACTCT	TAGACTTTTC	AAGTTGTTCT	CTTTGAAGCT	6780
TTTGAAAACC	TGCAATATCG	TTTGAATAGT	AAAGTGGGAT	AATCTGTGCC	ATCATAGACA	6840
CATGTTTCATG	ATTATGAAAA	TTCCTTGCCCT	TATCCATGAA	ATTTTCGATT	GTTACATGAA	6900
TGTTATCCAA	AATCTCAAAG	AAACGGGAGA	CTGCCAGGTC	AGACTCCCCA	AGCTCAAAGC	6960
GAGATAACTG	AGAGGTAGAG	CAGGATTCGC	CTGCTGCTTC	CTTTAAAGAA	TAATTTCCAC	7020
TTGTTTCGAAA	TTACAGAAAT	ACTTTTCCAA	GATGTTCCAT	CTTTACACCT	GCTCTGATAA	7080
TTCTTCCCAC	TCAAGCATAG	CTTCTTCCTG	ACGATGGCTG	ATTTTGTCCA	GCTCAGCCTG	7140
TAATTCATG	AGTTTGTCGG	CATCGTTTGT	TTCCAACATT	TGTTTCAGAAA	TGGCTTGGCT	7200
TTGACTTTCT	AGCTCTTCAA	TTTCAGCTTC	TAGACTTTTCG	ATTTGTGCGA	TGAGTTTGGC	7260
AACTTCTTTT	TGACTTTCTT	TCTGGGCCTG	ATAGTCATTG	ACTGGACTTG	CTTCCTTTGC	7320
TTGATTGCTA	GTTGAAGCTT	CCTCAGTCTG	ACTCATTTCT	GCTGTTGCTT	TCTTCTCAAC	7380
ATAGTAGTCG	TAATCTCCAA	GGTAGAGAGT	TGAACCATT	TCAGACAATT	CCAAAACATG	7440
AGTTGCCACA	CGATTGATAA	AGTAACGATC	ATGACTGACA	AACAGCAAGG	TTCCATCAAA	7500
GTCATCAAG	GCATTTTCTA	GCACTTCCTT	ACTATCAATA	TCCAAGTGGT	TGGTCGGCTC	7560

905

ATCCAGAATC	AAAAAGTTAT	TGTTTTCCAT	AGACAATTTA	GCTAAAAGCA	AACGAGCTTT	7620
TTCCGCCACCA	GATAGCATGC	CGACTGATTT	TTTAACATCA	TCTCCTGAGA	AAAGGAAGGC	7680
TCCAAGACGG	TTGCGGATTT	CAACTTCTGG	TGTCAGTTTG	AAATCATTCC	AGAGTTCATC	7740
CAGCACCGTA	TTACTTGGTG	TCAGCTTGCT	TTGGGTTTGG	TCATAGTAAC	CAACCTCAAC	7800
ATTAGCGCCA	AAGCGCTTTT	CTCCCTTGAT	AAAAGGAATC	TGGTCCACAA	TAGACTTGAT	7860
AAAGGTTGAC	TTGCCGATAC	CATTTGGACC	AACGATAGCG	ACAGCATTCA	TCTTACGAAG	7920
ATCTAGGTTA	ATCGGTTGTG	ACAAGACTTC	CCCCTCATAG	CCAACAGCTG	CATTTTCAAC	7980
AGTCAAAACA	ACATTGCCCG	ACGTTTTTTC	AGACTGGAAG	GTCATGTTGG	CTGATTTCTT	8040
GCCAGCTTCA	GGCTTGTC	AACGTTCCAT	TTTTTCCAGT	TGTTTACGGC	GAGATTGAGC	8100
ACGTTTAGTC	GTGAAGCAC	GAAGTAGATT	GCGATTGACA	AAGTCTTCCA	GAGCAGCGAT	8160
TTCCTTCTGT	TGCTTTTCAT	AGTTTTTTC	CTCAGTAACT	AGCTTTTGCT	CCTTCAATTC	8220
GACAAAACGA	GAGTAATTCC	CCACATAGCG	ATCCAAGGAA	TGCTTGGTCA	AATCTAGCGT	8280
AATTGTCGCA	ACCTTGTC	AGAAATAACG	GTCGTGGCTG	ACGATAATGA	GGGCACCGCT	8340
ATAGTTTACC	AAGTAATCT	CTAGCCAGGC	GATGGTTTCA	ATATCCAAGT	GGTTAGTTGG	8400
CTCGTCCAAG	ACCAAGAGAT	TGGGCTTTTC	AAGGAGCATT	TTGGCAAGTG	CCAAACGAGT	8460
ATTTTGACCA	CCAGAAAGCT	CAGCAATTTT	CATCTGCCAC	ATAGACTCGT	CAAACCTGAA	8520
TCCATTCAAA	ATCGCTCGAA	TATCAGCTTC	ATAGGTAAAG	CCACCTGCTT	GGCGAAAATT	8580
CTCAGATAAG	CGGTCATAAT	CTGACATCAG	TTTATCCAAA	TCCTCACCAG	ACTTTTCACC	8640
CATCTCCAGC	TCCATCTGAC	GCAGTTGTCT	CTCCGTCCGA	CGCAAATCAT	TAAAGACATG	8700
AAGCATTTCA	TCGTAGATGG	TATTTTCAGA	CTCAAACGG	CTATCTGGG	CTAGGTAAGA	8760
CAGAGAAATA	TCTTTTTTCT	TATGATTTT	TCCGCTAGTT	GGCTCCTCTT	CTCCAACATA	8820
AATCTTCAAA	AGAGTAGACT	TACCTGCACC	ATTTTCCCA	ACAAGAGCAA	TCCGATCTCG	8880
TTCATCAACC	TGCAGGTTGA	TATTATCGAA	AAGAACCTCT	CCTGCAAAAG	AACGTTCAAT	8940
TTTATTAGCT	TGTAAAATAA	TCATACAAGT	AGTATAGCAT	GTTTCCCTAA	GGCATTCAAG	9000
ATAATCGTAA	GTCTTTTAGT	ACAACCTTTA	TAACATAAAA	TAAACTAAAT	TATGTATATT	9060
TTATATTAGA	TTACTTCACT	ATCTTGTTGG	ATTTTCTAAC	CAGCTAATCT	TGTTTCAAAT	9120
AGTTATCGCA	CAAGTCTATT	ATTTAATCT	TTTCATCATT	TACGTACGTA	TAGCAGATTG	9180
AAATAAGATG	AGAACAAATC	GATTGGGAAA	GTAAAATTA	TTTCTATAAA	TGTTTTAGCA	9240
ATTGTTTCGT	ACTATTTTAG	ATTCAGTCTA	CTATATACAA	TATTTTCGGA	ACATTCAACT	9300

906

TTTAACTCT	ATTTATTACT	AGATTTTCATA	ATTAAAAAAC	CTACTGACCA	AGCTAGAAAG	9360
CTTGATACAA	TAGGCTTTTT	AAAGACTGAT	TATTTAACAG	CGTCTTTAAG	AGCTTTACCA	9420
GCTTTGAATG	CTGGTACTTT	AGAAGCTGCA	ATTGTCATTT	CTTTACCAGT	TTGTGGGTTG	9480
CGACCTTTAC	GTTCTGCGCG	CTCACGAACT	TCAAAGTTAC	CAAAACCGAT	CAATTGAACT	9540
T						9541

(2) INFORMATION FOR SEQ ID NO: 133:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 3502 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 133:

TTGACTATCC	TATCATGCTT	TCTAAGGTCT	ACTCAAGAAA	ATCATTTTCA	AGTTTTCACA	60
CCTTTCTCAA	AAAAGTTAAA	AAATTTTCTC	AAAAACGCTT	GACTCTGACC	TAAGGCGAAG	120
GTTATACTA	TCATTGTAAG	GAGGAAATCA	TGTACCATAT	AAAAGAAGCT	GCGCAGCTTT	180
CAGGTGTCTC	TGTC AAGACC	CTGCATCACT	ATGACAAGAT	AGGACTCTTG	GTCCCCTTAA	240
AGTCGGAAAA	CGGCTATCGA	ACCTACAGTC	AAGAGGATTT	GGAACGCCTT	CAGGTCATTC	300
TTTACTACAA	ATATCTAGGC	TTTTCTTTAG	AGAAAATAGC	AGAGCTGTTA	AAGGAAGAAA	360
GGACAGATTT	ATTGCCCCAT	TTGACTAGGC	AGTTGGACTA	TCTAACTCGC	GAAAGGCAAC	420
ATCTGGATAC	CTTGATTTCC	ACCTTGCAAA	AACTATTCA	AGAACA AAAA	GGAGAAAGAA	480
AAATGACCAT	TGAGGAAAAA	TTCACGGGAT	TTAGCTATCA	AGACAATCAA	AAATACCACC	540
AAGAAGCGGT	AGAGAAATAT	GGTCAAGAAG	TCATGGGACA	AGCGCTCGAA	CGCCAAAAAG	600
GTACGAAGA	CGAGGCTACG	GCCGCCTTTA	ACCAAGTCTT	TCAAAC TTTG	GCACAAAATC	660
TTCAAGTTGG	TTTACCTGCA	ACAGCAACCG	AAAACCAGGA	GCAAGCAGCC	AAGCTCTTGC	720
AAGCCATTCG	CACTTATGGA	TTTGACTGCT	CTATTGAGGT	ATTTCGGTCAT	ATCGGTAAAG	780
GTTACGTCTA	CAACCCAGAG	TTTAAGGAAA	ACATTGACAA	GTTTGGTTCT	GAAACAGCCC	840
AGTACACGTC	AGATGCCATT	GCGGTTTACG	TTCAGACAAA	TGCAGAATAA	ATAGGCTAGG	900
AATTTCCTAG	CCTATTTTTT	ACTTCAAATC	ATAAAGCCAG	TCGTCACCGT	TTTTGTAGTA	960
AAAGAATTCA	CTGAGATCTT	CTTCTAGAAA	CACACGAAGC	ATATCAGACA	TATCATCGGT	1020
TGCAAGTTTT	AGATGAGAAA	GATTTTCAAA	GTCCTCCAC	CAAAC TTTCC	CTTCGTCTGA	1080
AGACTGGAGT	TCACCAGTAA	AGTGTCTCTGT	CTTGTA AAAA	AGGACGACAT	AACGATAATC	1140

CTTGTCGTCA	TACCAGTTT	TGATACCACA	GAGTTGGGGT	TTGGAAATGA	TCAGACCAGT	1200
TTCTTCTTTC	ACTTCACGAA	TGACAGCATC	GACAAAGGAT	TCGCCACGTT	CAACATGACC	1260
ACCAGGAAAA	GTAATGCCAG	ACCAGTCGGG	ATTAACTCGG	TCTTGGACCA	GGACCTTATC	1320
TCCGTTTTTA	ATCATACACA	TGTAAACAAA	TTCGACTGCC	TCTCTTCTGT	TCATTCTTCA	1380
CAACCTTTAA	TCTTTAATCA	TAATGCAGAC	TTCCCGCCAC	CCAGCCGGTA	CAGAGGGCAG	1440
AAGTGATGTT	AAAGCCACCC	GTGTGGGCAT	TGATATCCAT	AACTTCGCCT	GCAAAGTGGA	1500
GGCCAGGTAC	CAGCTTACTT	TCAAGGGTTT	TAGGATTGAT	TTCTTGAGA	CTGACTCCAC	1560
CCTTGGTAAC	AAAGGACTTT	GCAAGGGACA	TTTTTCCAGT	TACAGGAATT	TTAAGTTCTT	1620
TAATGGACTG	GACAAGTTGT	TCTCGTTCCT	TTTCAGTCAG	TTGTTTACT	TTTTTCAGGAT	1680
ATCCTTGATC	AAAAAATTCG	GCCAAGCGTT	CTGGTAACAA	GGTTTTTAAA	GCGTTTTTCA	1740
AGGATTTTTC	CCGATTTTCT	TCTAGAAATG	TAACCAAGTC	CTTCTCAGAA	AGTTGAGGCA	1800
AAACATCGAG	TGAGAGAACC	TCCCCACCTT	TGACAAAGCT	AGACATGCGT	AGGGCAGCAG	1860
GACCTGACAA	ACCAAAGTGG	GTAAGAGTA	AATCATGAGT	GATGACATGC	TTACCATAAC	1920
TTAGGGTCAC	ATCGTCCAGA	GAAATACCTT	GTAAGGCTTT	ATGTGGAAAA	TCTGTTAATA	1980
AAGGACTTTC	AGCAGCCTCA	AGATCGGTGA	TGGTATGCTT	AAAATGGCGA	GCAATCTCGT	2040
GACCAAAACC	AGTCGAACCA	GTGGAAGGAT	AAGACTTACC	ACCTGTTGTG	ACAATGAGTT	2100
TCTCACAAGT	GAAGGTTTGA	TCCGCTGACT	TAAGGACAAA	CTGGTCATCT	ACTTTTTTAA	2160
CAGAAACGAT	TTCTATTTGA	GTAGCAACTT	GACCACCTAG	TTCCGGTATT	TTCTTTTCCA	2220
AAGCTTCGAT	AATAGTCCGA	GACTTGTAC	TGGCTGGAAA	GACGCGTCCG	TGGTCTTCGA	2280
CCTTAAGTTT	AACACCATTT	TCTGTAAAAA	AGTTGATGAT	GTCATGATTA	TCGAACTGGG	2340
AGAAAACACT	GTAAGAAAAG	CGTCCGTTTC	CAGGAATTCC	AGCTAGCAGG	TTGTCTAAGC	2400
TACCATTGTT	GGTCACATTG	CAACGTCCCC	CACCAGTCCC	AGCTAATTTT	TTTCCAAGTT	2460
TCCGATTTTT	TTCGATGAGG	AGGGTTTTCT	GTCCATAAAA	GCTACTGGAA	ATCGTAGCCA	2520
TCATACCAGC	AGGTCCCCCA	CCGATGACAA	TAGTATCAAA	ATGTTTCATA	GCTCTATTGT	2580
ACCACAAAAA	AACAAGAGAT	GATGGTCACC	TCTGTCAAG	AATGCAATTA	ATCAATTTCA	2640
TAGCCCATCA	GCAAACCGCC	CTCTCTGCA	TAGAAACTGC	AGAGACCAGA	GGTTGGTAGA	2700
ATTTTAATAT	CCGCTTGTGG	GAAGGTTTCA	CGGATTCGCT	CTGAGAGCTG	TTGACAACAT	2760
TTTTTCGTTAT	TGCGTTGGGC	CATGACAATA	CGGCCACCAG	CATATCCAGC	TTTTACTAAC	2820
TCATCATAGG	CAGCTTGAAC	TGATTTCTTT	GATCCCCTTG	CTTTTTGTAG	CAATTCGAGA	2880

908

GTCCCAGTTT CACTAGCTTT TCCGACCATA CGAATGTTGA GAAGGCCAAC GACCGTACCG	2940
ATAAGCTTGC TCAAACGGCC GTTCTTCACC AAGTTATCGA CTTTGGCTAG GACAAAGAGC	3000
AACTTAGTTT TTTCTTGATA GCGGGTGATA GCTTCAACCA CTTCTTCAA AGACAAGCCC	3060
TGGTCAATCA AGTCATTCAA TTTTCTACG AGTAGGTCAA CTTCAACCACC AGCAGATAAA	3120
CTATCAATCA CATGAATCTT AGTGTCAAGG TGCTCTTCCA GATAAATATT CTTTGCTAGT	3180
TGAGCACTAT TGTGACTGCC AGAAAGGGTA CCTGTGATGG TTAGTAGGAA AATGTTTTTG	3240
GCACCTTCAA ATGCTCGCAA ATAGTCATCT GGGCTTGGAC AAGCCGATTT TGAAGCTTCT	3300
GCAGTTGCAT ACATGGTTTC CATCATTTGG TCAATATCGA GACTGGCGTC ATCAACAAAG	3360
ACCTGATCAG CTACTTGAAT GGTAAAGGGG ACACCTACAA AGGTTGTGTT AATAGCTGGT	3420
GTTGGCAGTT GACGATAATC ACAACCAGAG TCAGCAATAA TCTTCCAAGT CATAGAAATT	3480
CTCCATCTTT GTCAGGAACG AT	3502

(2) INFORMATION FOR SEQ ID NO: 134:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 12665 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 134:

CGATTGATTT TTTTAAAGCG TTCGATAGAG AATGAGAAAC GAATCCTTAG CAATGGCGGG	60
AAAGAATTTG GAGTTGAGAA TACAAAACGA TTAACATATGG CTCATATTGT TTTTATCTC	120
TCTTGCTTGG TTGAGGCAAT GGTGCACAAG ACAATTTTGG ATGGCATGGG CATGGTTGGT	180
TTAGTCTTGC TTATTTTTTC TATGCTGATG TTGATGTTGG TGATTCACTT GTTGGGAGAT	240
ATTTGGACAG TGAAGCTTAT GCTTGTCAAT AATCACAAAT ATGTAGATCA TATCTTGTTT	300
AGGACAGTAA AACACCCTAA TTAATTTTTA AATATCTTTC CTGAGTTGAT TGGCTTGACC	360
TTGTTGAGTC ATGCTTATGT GACTTTTGTT TTAGTTTTTC CAGTTTATGC AGTTATTTG	420
TATCGACGAA TAGCTGAAGA GGAAAAGCTA TTACATGAAG TTATAATCCC AAATGGAAGC	480
ATAAAGAGAT AAATACAAAA TTCGATTTAT ATACAGTTCA TATTGAAGTG ATATAGTAAG	540
GTTAAAGAAA AAATATAGAA GGAATAAAC ATGTTTGCAT CAAAAGCGA AAGAAAAGTA	600
CATTATTCAA TTCGTAAATT TAGTGTTGGA GTAGCTAGTG TAGTTGTTGC CAGTCTTGT	660
ATGGGAAGTG TGGTTCATGC GACAGAGAAC GAGGGAGCTA CCCAAGTACC CACTTCTTCT	720
AATAGGGCAA ATGAAAGTCA GGCAGAACAA GGAGAACAAC CTAAAAAATCGATTTCAGAA	780

909

CGAGATAAGG CAAGGAAAGA GGTCGAGGAA TATGTAAAAA AAATAGTGGG TGAGAGCTAT	840
GCAAAATCAA CTAAAAAGCG ACATACAATF ACTGTAGCTC TAGTTAACGA GTTGAACAAC	900
ATTAAGAACG AGTATTTGAA TAAAATAGTT GAATCAACCT CAGAAAGCCA ACTACAGATA	960
CTGATGATGG AGAGTCGATC AAAAGTAGAT GAAGCTGTGT CTAAGTTTGA AAAGGACTCA	1020
TCTTCTTCGT CAAGTTCAGA CTCTTCCACT AAACCGGAAG CTTTCAGATAC AGCGAAGCCA	1080
AACAAGCCGA CAGAACCAGG AGAAAAGGTA GCAGAAGCTA AGAAGAAGGT TGAAGAAGCT	1140
GAGAAAAAAG CCAAGGATCA AAAAGAAGAA GATCGTCGTA ACTACCCAAC CATTACTTAC	1200
AAAACGCTTG AACTTGAAAT TGCTGAGTCC GATGTGGAAG TTAAAAAAGC GGAGCTTGAA	1260
CTAGTAAAAG TGAAAGCTAA CGAACCTCGA GACGAGCAA AAATTAAGCA AGCAGAAGCG	1320
GAAGTTGAGA G'AAAACAAGC TGAGGCTACA AGGTTAAAAA AAATCAAGAC AGATCGTGAA	1380
GAAGCAGAAG AAGAAGCTAA ACGAAGAGCA GATGCTAAAG AGCAAGGTAA ACCAAAGGGG	1440
CGGGCAAAAC GAGGAGTTCC TGGAGAGCTA GCAACACCTG ATAAAAAAGA AAATGATGCG	1500
AAGTCTTCAG ATTCTAGCGT AGGTGAAGAA ACTCTTCCAA GCCCATCCCT GAAACCAGAA	1560
AAAAGGTAG CAGAAGCTGA GAAGAAGGTT GAAGAAGCTA AGAAAAAGC CGAGGATCAA	1620
AAAGAAGAAG ATCGCCGTAA CTACCCAACC AATACTTACA AAACGCTTGA ACTTGAAATF	1680
GCTGAGTCCG ATGTGGAAGT TAAAAAAGCG GAGCTTGAAC TAGTAAAAGA GGAAGCTAAG	1740
GAACCTCGAA ACGAGGAAAA AGTTAAGCAA GCAAAAGCGG AAGTTGAGAG TAAAAAAGCT	1800
GAGGCTACAA GGTTAGAAAA AATCAAGACA GATCGTAAAA AAGCAGAAGA AGAAGCTAAA	1860
CGAAAAGCAG CAGAAGAAGA TAAAGTTAAA GAAAAACCAG CTGAACAACC ACAACCAGCG	1920
CCGGCTCCAA AAGCAGAAAA ACCAGCTCCA GCTCCAAAAC CAGAGAATCC AGCTGAACAA	1980
CCAAAAGCAG AAAAACCAGC TGATCAACAA GCTGAAGAAG ACTATGCTCG TAGATCAGAA	2040
GAAGAATATA ATCGCTTGAC TCAACAGCAA CCGCCAAAAA CTGAAAAACC AGCACAACCA	2100
TCTACTCCAA AAACAGGCTG GAAACAAGAA AACGGTATGT GG'FACTTCTA CAATACTGAT	2160
GGTTCAATGG CGACAGGATG GCTCCAAAAC AATGGCTCAT GG'ACTACCT CAACAGCAAT	2220
GGCGCTATGG CGACAGGATG GCTCCAAAAC AATGGTTCAT GG'ACTATCT AAACGCTAAT	2280
GGTTCAATGG CAACAGGATG GCTCCAAAAC AATGGTTCAT GG'ACTACCT AAACGCTAAT	2340
GGTTCAATGG CGACAGGATG GCTCCAATAC AATGGCTCAT GG'ACTACCT AAACGCTAAT	2400
GGTTCAATGG CGACAGGATG GCTCCAATAC AATGGCTCAT GG'ACTACCT AAACGCTAAT	2460
GGTGATATGG CGACAGGTTG GGTGAAAGAT GGAGATACCT GG'ACTATCT TGAAGCATCA	2520

GGTGCTATGA AAGCAAGCCA ATGGTTCAAA GTATCAGATA AATGGTACTA TGTC AATGGC 2580
 TCAGGTGCC TTGCAGTCAA CACAACGTGA GATGGCTATG GAGTCAATGC CAATGGTGAA 2640
 TGGGTAAACT AAACCTAATA TAACTAGTTA ATACTGACTT CCTGTAAGAA CTCTTTAAAG 2700
 TATTCCCTAC AAATACCATA TCCTTTCAGT AGATAATATA CCCTTG TAGG AAGTTTAGAT 2760
 TAAAAAATAA CTCTGTAATC TCTAGCCGGA TTTATAGCGC TAGAGACTAC GGAGTTTTTT 2820
 TGATGAGGAA AGAATGGCGG CATTCAAGAG GCTCTTTAAG AGAGTTACGG GTTTTAAACT 2880
 ATTAAGCCTT CTCCAATGTC AAGAGGGTTT CAATCTCTGC CAGGGTGCTG GCTTGCGAAA 2940
 TGGCTCCACG GAGTTTGCA GCGCCAGATG TTCCACGGAG ATAGTGAGGA GCGAGACCGC 3000
 GGAATTCACG AACTGCGACG TTTTCTCCTT TGAGGTAAAT CAATCGTTTC AAGTGTTCGT 3060
 AGGCGATCTT CATCTGTCT TCAAAGGTCA AATCAGGTAG GATTTCTCCT GTTCAAAGT 3120
 AATGGTTGAT TTGGTTGAAG AGGTAAGGAT TTCCCATGGC AGCTCGGCCA ATCATGACTG 3180
 CGTCAGCAC AACTTCTTCG ATGCGTTGCT TGGCTTCTTG GACAGTACGG ATATCACCGT 3240
 TGGCGATGAA TGAATCTTG GTTAGAGCTT GGGCAACCTT GTAAAGGGTC TCAAGGTCTG 3300
 CGTGCCAGT ATACATTTGT TCACGGGTAC GGCCATGCAT GGCGAGGGCA GAAACACCTG 3360
 CAGCTTCAGC AGCGAGAGCA TTTTCTACTG CAAGAGATGG GTCCGCCAG CCGGTACGCA 3420
 TTTTGACAGT AAGTGGGATA TCAAGGACAG ACTGGACCTT GTTGATGATG GAGTAAATCT 3480
 TGTCTGGATC CTFGAGCCAC ATAGCACCAG CTTCTCTT CACGATTTTG TTGACAGGGC 3540
 AGCCCATGTT GATATCGACG ATATCGGTCT TGGTGTTC TTGGATGAAT TCTGCTGCGC 3600
 GTGCTAGGCT GTCTTCATCG CTACCAAAAA GTTGGATAGA GACAGGGTT TCGCCCTCAT 3660
 CGATATGAAG CATGTGCAGG GTTTTTTCGT TGTGTATTG GATTCCCTTG TCAGAGACCA 3720
 TTTCCATTAC AACGAGTCCA GCTCCGAGCT CCTTTGCGAT AGTACGAAAG GCTGAGTTGG 3780
 TCACGCCAGC CATAGGCGCT AAAACGGTAC GATTGGGAAT CTCAATATG CCAATCATAA 3840
 AAGGTGTATT AAGATTTGTC ACGAATGAGT TCCTCCAGGT CCTTTTCATC AAAGTTGTAA 3900
 GTAGTTTGGC AGAATTGACA AGTGATTTCT GCCCCGTGGT CTTCCTCTTT CATTTCTGT 3960
 AAGTCTGAGC TTGGAAGGCT GGCAAGAGCG TTCATAAAGC GTTCATGGCT ACAGTCACAT 4020
 TGGAAACGGA TTTCTTCTTC AGAAAGACGC TTGTAGGCTT CGTCCCCGTA GATAGCCTTG 4080
 AGGAGGGCTT CGATATGGTC GTCGCTTTCG AGAAGAGTAG AGATAGCTGG CATTTCTTGG 4140
 ATGCGTTTTT CAAAGCGAGC AATCTCTTCT TTCTTGCTC CTGGCAAGAC TTGAACTAGG 4200
 AAACCACCTG CAACCTTGAC CTTGTCTTCC TCGTCCAAA GGACATGAG GCCGACCGCT 4260
 GAAGCGTTT GTTGGCTTTC AGTAAGGTAA AAGGCAAGGT CTTACCGAT TTCTCCAGAG 4320

ATGAGGGGAG TTATAGAGTT GTAAGGATTT CCAGTACCGT AGTCTGTGAT AACGAGGAAT 4380
TGACCATTTT CAACAAAAGG TCCGACTAGG ACTTCACCAG TCGCAGTCTT TTTGATGTCA 4440
ACACCAGGAT TTTGAACATA GCCTTTGACG TTCCCCTTGG TATCAGCGAC GGTGATAATA 4500
GCACCTAGAG AGCTAGATCC CAACACCTTA ACTGTAAGTT TGGTATTTCC TTTTTCATFG 4560
GCTGCGAGAA TCTGGCTAGC GATAAGAGTT CGACCAAGCG CTACAGTTGA GCTAGCTTGG 4620
GTTTGTATGTT TTTCTTGAGC AGTGCGGACG GTTTCAGTGC TATCAAGGAC AAAAGCACGA 4680
AAGGcTCCGC TTTCTGATAT AGTTTTAATA ATTTTATCCA TAGCTACTAT TTTAGCATAA 4740
AAATGCCCAA AGGGGGAGCC GTGTGTTTAC TGATTTTCAG GATAATGGAC CAGGAAATCA 4800
GCATGAAAAAT AAAAAGAGAA ACAGATTATT TTAGCATTTG TCAGATTTAT GCTATGCTTA 4860
AGGTAGAAAA TGAAGGGAT AACAAATGTA TTTAGGAGAT TTGATGGAGA AAGCCGAGTG 4920
TGGTCAATTT TCAATACTTT CCTTCTATT ACAAGAGTCT CAGACGACCG TCAAGGCTGT 4980
AATGGAAGAA ACAGGATTTT CAAAAGCAAC CCTAACCAA TATGTCACCC TGCTCAATGA 5040
CAAGGCTTTG GATAGTGGCT TAGAGCTGGC TATTCACTCA GAAGATGAAA ATCTGCGTCT 5100
GTCTATCGGT GCAGCTACCA AGGGGAGAGA TATTCGGAGC TTGTTTTTGG AGAGTGCTGT 5160
TAAATACCAG ATTTTGGTTT ATCTTCTCTA CCACCAACAG TTTTLAGCCC ATCAGCTGGC 5220
TCAAGAATTG GTGATTAGCG AGGCTACGCT TGGTCGTCAC TTGGCTGGTT TAAATCAGAT 5280
TTTGTGAGAA TTTGATTAT CCATCCAAA TGGCCGTTGG CGAGGTCCAG AGCATCAGAT 5340
TCACTATTTT TATTTCTGTC TTTTCCGAAA GGTCTGGTCG AGTCAGGAAT GGAAGGTCA 5400
CATGCAGAAA CCAGAGAGAA AACAGGAGAT TGCCAATTTA GAGGAAATCT GCGGTGCAAG 5460
TTTGTCTGCG GGGCAGAAAT TGGACTTGGT TCTCTGGGCT CACATCAGTC AACACGTCT 5520
TCGGGTCAAT GCTTGTGAGT TTCAAGTCAT AGAAGAGAAA ATGCGAGGGT ATTTTGACAA 5580
TATCTTTTAT CTTCGTTTGC TGAGAAAGGT TCCGTCCTTT TTTGCTGGGC AACATATTC 5640
ACTAGGAGTT GAGGATGGTG AGATGATGAT ATTCTTCTCT TTTCTCCTAT CTCATCGCAT 5700
TCTTCTCTT CATACTATGG AGTATATCTT TGGTTTTGGA GGGCAGTTGG CAGATTTACT 5760
GACGCAATTG ATTCAAGAAA TGAAGAAGGA GGAACATTTG GGGGATTATA CAGAGGACCA 5820
TGTCACCTAT GAACTCAGTC AGCTTTGTGC TCAAGTCTAT CTCTATAAGG GCTATATTTT 5880
ACAGGATCGC TACAAGTACC AGTTAGAGAA TCGTCATCCA TATTTACTGA TGAACATGA 5940
TTTTAAAGAG ACAGCAGAGG AGATTTTCA TGCTCTACCT GCTTTTCAAC AGGGGACAGA 6000
TTTAGATAAG AAGATTCTCT GGAATGGCT CCAGTTAATC GAATATATGG CTGAAAACGG 6060

912

TGGCCAGCAT	ATGCGGATTG	GTCTGGATTT	GACATCTGGT	TTCCTTGTCT	TTCAAGGAT	6120
GCCAGCCATT	TTGAAACGGT	ATTTGGAATA	CAATCGTTTT	ATTACCATTG	AAGCTTATGA	6180
CCCTAGTCGG	CATTATGATT	TGCTGGTTAC	CAATAACCCG	ATTCATAAGA	AGGAACAGAC	6240
ACCAGTCTAT	TATTTAAAAA	ATGACTTGGA	TATGGAGGAT	TTGGTAGCGA	TTGCCAGTT	6300
ATTATTCACT	TAAAAGGCTT	GGTTAATCCA	GGTCTTTTTT	GTGAAATCA	CACAATCTCC	6360
TCACATTTTT	TTAAAAATTA	AAAAAAGTTG	ATAAACAAAG	AAGCGCTTTA	TTTTGTATAC	6420
TAGTAAGTGT	AAAGAGGAAA	CACCTCAAGA	TCTTTATCAG	GAGGACAGTA	CATGTCACAA	6480
GAAAAATACA	TCATGGCCAT	TGACCAGGGA	ACTACAAGTT	CTCGTGCCAT	CATTTTCAAC	6540
AAAAAAGGGG	AAAAGGTTAG	CTCGAGTCAA	AAAGAGTTTA	CCCAGATTTT	CCCTCAGGCA	6600
GGTTGGGTTG	AGCACAAATG	CAATGAAATT	TGGAACTCTG	TTCAGTCAGT	TATTGCGGGT	6660
GCTTTCATCG	AAAGTGGTGT	CAAGCCAAAT	CAAATCGAGG	CAATCGGGAT	TACCAACCAA	6720
CGTGAACAA	CGGTGTCTG	GGATAAGAAA	ACAGGACTTC	CTATCTACAA	TGCTATCGTT	6780
TGGCAGTCAC	GCCAGACAGC	ACCTTTGGCT	GAGCAACTAA	AAAGCCAAGG	TTATGTGGAA	6840
AAATTCACAT	AAAAGACTGG	TTTGATTATT	GATGCTTACT	TCTCTGCTAC	CAAGGTFCGT	6900
TGGATTTTGG	ATCATGTAGA	AGGTGCTCAA	GAGCGAGCAG	AAAAAGGGGA	ATTGCTCTTT	6960
GGTACTATCG	ATACTTGGTT	GGTTTGAAA	TTGACTGACG	GTGCGGCTCA	CGTGACTGAC	7020
TACTCAAATG	CAGCTCGTAC	CATGCTTTAT	AACATTAAAG	AACTCAAATG	GGATGATGAG	7080
ATTTTGAAA	TCCTTAACAT	TCCGAAGGCT	ATACTTCCAG	AAGTTCGTTT	TAACCTCCGAA	7140
ATCTACGGCA	AGACAGCTCC	ATTCCATTTT	TACGGTGGAG	AGGTGCCAAT	CTCAGGTATG	7200
GCTGGGGACC	AACAAGCAGC	CCTCTTTGGA	CAGTTGGCTT	TTGAGCCAGG	TATGGTTAAG	7260
AATACTTATG	GAACAGGCTC	TTTCATCATC	ATGAATACTG	GGGAAGAGAT	GCAGTTGTCT	7320
GAAAACAACC	TCTTGACAAC	CATTGGTTAC	GGAATCAACG	GTAAGGTTTA	TTATGCCTTG	7380
GAAGGTCTA	TCTTCATCGC	AGGAAGTGCT	ATTCAGTGGC	TTCGTGACGG	TCTTCGCATG	7440
GTTGAAAATT	CACCAGAATC	TGAAAAATAC	GCTCGTGATT	CTCACAACAA	CGATGAAGTT	7500
TATGTCGTTT	CAGCCTTTAC	AGGTCTAGGC	GCTCCATACT	GGAACCAAAA	TGCTCGTGGT	7560
TCCGTCTTTG	GTTTGACTCG	TGGAACAAGC	AAAGAAGACT	TTATCAAGGC	GACTTTGCAA	7620
TCTATGCTT	ATCAAGTGCG	TGATATCATC	GACACCATGC	AAGTGGATAC	TCAGACCGCC	7680
ATTCAAGTAC	TGAAGGTGGA	TGGTGGTGCA	GCCATGAACA	ACTTCCTCAT	GCAGTTCCAG	7740
GCGGATATTT	TAGGCATTGA	CATTGCACGT	GCTAAAAACC	TGGAAACAAC	AGCTCTAGGA	7800
GCGGCCTTCC	TAGCAGGTTT	GTCAGTAGGG	TACTGGAAAG	ACTTGGACGA	GTTGAAACTC	7860

TTGAACGAGA	CAGGAGAACT	CTTTGAGCCA	TCTATGAACG	AATCTCGCAA	GGAACAACCTC	7920
TACAAGGGCT	GGAAGAAGGC	TGTGAAAGCA	ACTCAAGTCT	TTGCGGAAGT	AGACGACTAA	7980
TACTGGCAGA	ATAAAGCGAT	TTATTTAGAA	AGTGTGTAAG	TATGGAATTT	TCAAAGAAAA	8040
CACGTGAATT	GTCAATTAAA	AAAATGCAGG	AACGTACCCT	GGACCTCTTG	ATTATCGGTG	8100
GAGGAATCAC	AGGAGCTGGT	GTAGCCTTGC	AGGCGGCAGC	TAGCGGTCTT	GAGACTGGTT	8160
TGATTGAAAT	GCAAGACTTT	GCAGAAGGAA	CATCTAGTCG	TTCAACAAAA	TTGGTTCACG	8220
GAGGACTTCG	TTACCTCAAA	CAATTTGACG	TAGAAGTGGT	CTCAGATACG	GTTTCTGAAC	8280
GTGCAGTGGT	TCAACAAATC	GCTCCACACA	TTCCAAATC	AGATCCAATG	CTCTTACCAG	8340
TTTACGATGA	AGATGGAGCA	ACCTTTAGCC	TCTTCCGTCT	TAAAGTAGCC	ATGGACTTGT	8400
ACGACCTCTT	GGCAGGTGTT	AGCAACACAC	CAGCTGCGAA	CAAGGTTTTG	AGCAAGGATC	8460
AAGTCTTGGA	ACGCCAGCCA	AAC TTGAAGA	AGGAAGGCTT	GGTAGGAGGT	GGAGTGATC	8520
TTGACTTCCG	TAACAACGAT	GCGCGTCTCG	TGATTGAAAA	CATCAAACGT	GCCAACCAAG	8580
ACGGTGCCCT	CATTGCCAAC	CACGTGAAGG	CAGAAGGCTT	CCTCTTTGAC	GAAAGTGGCA	8640
AGATTACAGG	TGTTGTAGCT	CGTGATCTCT	TGACAGACCA	AGTGTTTGAA	ATCAAGGCCC	8700
GTCTGGTTAT	TAATACAACA	GGTCTTGGA	GTGATAAAGT	ACGTAATTTG	TCTAATAAGG	8760
GAACGCAATT	CTCACAAATG	CGCCCAACTA	AGGGAGTTCA	CTTGGTAGTA	GATTC AAGCA	8820
AAATCAAGGT	TTCACAGCCA	GTTTACTTCG	ACACAGGTTT	GGGTGACGGT	CGTATGGTCT	8880
TTGTTCTCCC	ACGTGAAAAC	AAGACTTACT	TTGGTACAAC	TGATACAGAC	TACACAGGTG	8940
ATTTGGAGCA	TCCAAAAGTA	ACTCAAGAAG	ATGTAGATTA	TCTACTTGGC	ATTGTCAACA	9000
ACCGTCTCCC	AGAATCCAAC	ATCACCATTG	ATGATATCGA	AAGCAGCTGG	GCAGGTCTTC	9060
GTCCATTGAT	TGCAGGGAAC	AGTGCCTCTG	ACTATAATGG	TGGAAATAAC	GGTACCATCA	9120
GTGATGAAAG	CTTTGACAAC	TTGATTGCCA	CTGTTGAATC	TTATCTCTCC	AAAGAAAAAA	9180
CACGTGAAGA	TGTTGAGTCT	GCTGTCAGCA	AGCTTGAAAG	TAGCACATCT	GAGAAACATT	9240
TGGATCCATC	TGCAGTTTCT	CGTGGGTCTA	GCTTGGACCG	TGATGACAAT	GGTCTCTTGA	9300
CTCTTGCTGG	TGGTAAAATC	ACAGACTACC	GTAAGATGGC	TGAAGGAGCT	ATGGAGCGCG	9360
TGGTTGACAT	CCTCAAAGCA	GAATTTGACC	GTAGCTTTAA	ATTGATCAAT	TCTAAAACCT	9420
ACCCTGTTTC	AGGTGGAGAA	TTGAACCCAG	CAAATGTGGA	TTCAGAAATC	GAAGCCTTTG	9480
CGCAACTTGG	AGTATCACGT	GGTTTGGATA	GCAAGGAAGC	TCACTATCTG	GCAAATCTTT	9540
ACGGTTCAAA	TGCACCGAAA	GTCTTTGCAC	TTGCTCACAG	CTTGAACAA	GCGCCAGGAC	9600

914

TCAGCTTGGC	AGATACTTTG	TCCCTTCACT	ATGCAATGCG	CAATGAGTTG	ACTCTTAGCC	9660
CAGTTGACTT	CCTTCTTCGT	CGTACCAATC	ACATGCTCTT	TATGCGTGAT	AGCTTGGATA	9720
GTATCGTTGA	GCCAATTTTG	GATGAAATGG	GACGATTCTA	TGACTGGACA	GAAGAAGAAA	9780
AAGCAACTTA	CCGTGCTGAT	GTGGAAGCAG	CTCTCGCTAA	CAACGATTTA	GCAGAATTAA	9840
AAAATTAAGA	AAAAATAAAA	GAGGTGGAGG	GCAGCATTCC	TTGTCGCCCG	TCCCTTCTTT	9900
TTAATGGAGA	CAGAAAGATG	ATGAATGAAT	TATTTGGAGA	ATTTCTAGGG	ACTTTAATCC	9960
TGATTCTTCT	AGGAAATGGT	GTTGTTGCAG	GTGTGGTTCT	TCCTAAAACC	AAGAGCAATA	10020
GCTCAGGTTG	GATTGTGATT	ACTATGGGTT	GGGGGATGTC	AGTTGCCGGT	GCAGTCTTTG	10080
TATCTGGCAA	GCTCAGTCCA	GCTTATTTAA	ACCCAGCTGT	GACCATCGGT	GTGGCCTTAA	10140
AAGGTGGTTT	GCCTTGGGCT	TCCGTTTTGC	CTTATATCTT	AGCCAGTTC	GCAGGGGCCA	10200
TGCTGGGTCA	GATTTTGGTT	TGGTTGCAAT	TCAAACCTCA	CTATGAGGCA	GAAGAAAATG	10260
CAGGCAATAT	CCTGGCAACC	TTCAGTACTG	GACCAGCCAT	CAAGGATACT	GTATCAAACCT	10320
TGATTAGCGA	AATCCTTGGA	ACTTTTGTTT	TGGTGTGAC	AATCTTTGCT	TTGGGTCTTT	10380
ACGACTTTCA	GGCAGGTATC	GGAACCTTTG	CAGTGGGAAC	TTTGATTGTC	GGTATCGGTC	10440
TATCACTAGG	TGGACAACA	GGTTATGCCT	TGAACCCAGC	TCGTGACCTT	GGACCTCGTA	10500
TCATGCACAG	CATCTTGCCA	ATTCCAAACA	AGGGAGACGG	AGACTGGTCT	TACGCTTGGA	10560
TTCTGTGTG	AGGCCCTGTT	ATCGGAGCAG	CCTTGGCAGT	GCTTGTATTC	TCACTTTTCT	10620
AGTTTATACT	CTTCGAAAAT	CAAATTCAAA	CCACGTCAGC	GTCGCCTTAC	CGTACTCAAG	10680
TACAGCTTGC	GGCTAGCTTC	CTAGTTTGCT	CCTTGATTTT	CATTGAGTAT	TAGAAAACAA	10740
TTATGTTGAT	AGAGCTTGGG	CAAGAGCCCA	ATTTAGCAA	AAAATGAAGT	AAATCTTCTC	10800
ATAATAAAAC	GCATCATATC	AAGCACGAAA	ATCCACGAG	GTCAACTACA	GTCAGAAAGC	10860
TGAACAACAA	GCCAAAACGC	CCAAAAAAGG	CGGCAAAAAG	CAAGCACCTG	CAAGCAACGT	10920
GCCGAAATGG	TCAAATCCTG	ATTATGTCAA	CGAATTAGAC	CCAAAAATCG	TTGATATGCT	10980
AGTAGAATTT	CACAAGTCAC	AAGGCACTTT	GGAAACTCCC	GAGGCGCAAG	CAGAAATCCG	11040
CCAAAAACGT	GAAGAAATCG	AGCAAAGGAG	AGCTGAGCTT	GAGGGTAAAA	AACAAGAGCT	11100
TTTGAACCGC	TTGAACAAAT	AGAGTTTCGC	AAGTATTATG	CTTACAAATT	ACTTGAGCAA	11160
TTAACTAAAA	TATAAACCTT	GCCTTTATAT	CTAGGCAGGG	TTTATATTTT	AGAAATTCAC	11220
GTAGGTTGTT	ACGGTTTTTA	CATACCCAGT	ATAGTTTGAG	TTTCTATAGT	ATTCAGTGAT	11280
AAACTTCCAT	TTTCTTTGAG	CAACATGGAT	ATAAGTACTT	GTTATGTAGT	ATGGATATGG	11340
GCTTTGTGAA	TCCAAGTAAG	ACTGATAAGC	TTGTATACCA	AAATATGCTC	CACCAATTAT	11400

915

TGCACCCCAT GGACCCCCA ATAAAGCACC TATCCTACCA ATCATATAAC TGATTCCAGC 11460
 ACCAGTCATG AAGTTAGCGA ATGTGTTAGC TTGTTTATTC CCATGTATTG TGTGACGTA 11520
 ATTCCAAACA TTAGGATCGT ATGATCTAAA AGATATATTT AGGTCGATTT CATTCCTTTG 11580
 ATAAGCCATA TAAAATGCCC CATTGATATA GACGCCGTCA GCACGTCGTT CAATAGTGTC 11640
 TACTTCCA TCTGGATTGA CAACCTCAAG AACTTCATCG CTAAAATAT TTAATTGCGT 11700
 ATCTCCGAAC CGCACTGATG AGCCATTCTC AACTGAGCC TCACCAGATA CAACTTTAGA 11760
 GTTTGCCGAT AAGCTATCAT CAGCAAAAAC AAACAAGCGA CGGGGAAATG CTAGACATA 11820
 AGAAAACAGA CATAACTAGC AAACACATGC ATTTAAACAT CTTAGACATA ACGGAAACTC 11880
 CTTTGTATTT TTGATTTTTT TCAACTTTTA TTATACAATA AAACCAAATA AAAAGAAAGC 11940
 GGTAACAATA TGCTTAATGC GAAAATTTTT TATATATTTT TATGTTTGGT CGTTATCGAA 12000
 ACTACAGGCT TGTGTTGTT GAAAAGAGGT CTCGAAATGG GTTATTTAGA CACAGAAGCT 12060
 ATTATCCTCG CAGTTTTTTC ATTTGCTTFT TACAACCTAT GTTCATTGCG TTGGGTCTGC 12120
 TCTACAATAA AAAACAATA AAAATAATA GACGTATTTT CAAAAAAAC maAATGCATA 12180
 TTTATATTAG CAAAACGACG ATTTAAATCG TCGTTTTTTT GTAGTACGAC GGGCATGTGCG 12240
 TATATCTGAG GTGTAAGTCC TCAGCCTGAC TATCGTGAGG TAGCAGGGAG AGGAAGGGAT 12300
 AGCGAAATCG TGGCTCTACG AACAGGAACG TGATAGTAAG GCGTATATAG CGGATAAGGA 12360
 GGCTTCAAAC TCTAAAGTCC AAAAAGGTAG TCGTAACCTA TATGTGTAAA TCACGAGAGT 12420
 AATTGAATTC GGACTAAGGT TTGTGTGAAA AAGATAAATC TTTCTAGAGT CTAAAGACTC 12480
 TCGGTCAGAT TTCCTATTTT CACTGTAACC TTTAACGTC CTCATATCTT GTATAACGA 12540
 GGAAAGATGT ACGACTTATC CCGTGAGGTT TCATGAGCGT GAAAGCGTAG TAACAACGAA 12600
 TCATGAGAAG TCAGCCGAGC CCATAGTAGT GAGGAACTT CCGTAATGGA AGTGGAGCGA 12660
 AGGGG 12665

(2) INFORMATION FOR SEQ ID NO: 135:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 5305 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 135:

CGCTAATCAC TACAATCATT TTATTGTACT TTTTCACTCT CAAGAAAAGC AAGAAGTATT 60

CATTTTAGTT	TCATTTAGTA	TTATTTTGCA	TACCTAAAAT	ACAGTAAAAA	ATCAGTCATC	120
TTGGTATGCT	CCTGCTTTCA	CTATTCACA	CGTTTTTGAC	TTATACTAGG	CTCATTTCCA	180
AAAGCATTAT	ATAATAGTGA	TATGAAACCA	ACTAAACTAA	ACAAGAAATA	TAAGCAATAA	240
AAATTCGTTT	AAAAGATCTT	ACTAAAGCTA	ATACTAAATA	AAAATAAAAAG	AGTAAACTAG	300
GAAGTTTATT	TCAAACAACC	TAAAATACTG	ATTTTCGGCT	GAAGATAATA	CTGGAGTGCA	360
AATTAATGGG	GTTATAATAA	ATAGCTGATA	GCTTGTGTTG	GTTTTGGATT	TTTTAAGAGT	420
AGATGAGTAT	TAAAACTATA	AGGAGGACGA	AGGTGGCTAA	AAATTTAAAA	TTAAAATTAG	480
CTCGGGTAGA	GCGTGATTTA	ACACAAGGTC	AACTGGCAGA	GGCTGTCGGG	GTGACACGCC	540
AGACTATTGG	TTTAATAGAG	GCGGGAAAAT	ACAATCCCAG	TCTCTCGCTC	TGCCAGTCTA	600
TTTGCAGATG	TTTAGGGAAA	ACCCTAGACC	AACTATTTTG	GGAGGAAGAA	GATGAAAAAT	660
AGATTTTATT	ATTCTCAATT	ACTAGACGAA	AGAGAAGAAC	AACTGTTCAA	TAAAGCGGGC	720
TCTGAAAGTT	TCTATATCTG	CATTGCTTTG	TCGCTCCTAT	CTTATATCAT	TTCAGTATTA	780
GCACCAAGCC	TTTTTAATTC	TAATATGCTG	CTAATCGTTA	TCATCATAGG	GACATTTTAC	840
TTTTTCAATC	GTGCCCGTTA	TCTGGGAGTG	ACCTACTATG	GTCGTTTTCA	TTTTACGATT	900
TTGGGTGTTT	TTTTCCTAAC	CTTGGCTATT	ACGGCTCTTT	TGATGTGCA	GAATTATCAA	960
TTCAACATAG	AAATTTATCA	GCACAATCCT	TTGAATTTTA	AATACCTGTC	TGCTTGGGTC	1020
ATTACTTATA	TCATTTACCT	TCCGTGGATC	TTTATTGGCA	ATCTTGGTCT	TAAGAGCTAT	1080
GGCGAATGGG	CTCAGAAAAA	ATTTGAACAA	GATATGGATG	AATTGGAGAG	TGGAGAATAG	1140
CTTGTTACTC	TTTTCTCAAT	CCAGCTAAAA	TGTGATATAA	TAGTACTAAT	TTATTGGAAT	1200
ACATGAAAGT	TCTTGAAAAA	TTTCATGGGT	TTCTAGCTAA	GGAAGTAGGA	AAAGTATGTA	1260
TCCAGATGAT	AGTTTGACAT	TGCACACGGA	CTTGTACCAG	ATCAACATGA	TGCAGGTTTA	1320
CTTTGACCAA	GGGATTCACA	ATAAGAAGGC	GGTCTTTGAG	GTGTATTTCC	GCCAACAGCC	1380
TTTTAAGAAC	GGCTATGCGG	TTTTTGACAG	TTTAGAAAAGA	ATTGTGAACT	ATCTTGAAGA	1440
CTTGCGTTTT	TCAGATAGTG	ATATAGCCTA	TTTGGAGTCG	CTTGGTTATC	ATGGGGCGTT	1500
CTTGGATTAC	CTTCGCAATT	TCAAGTTGGA	GTTGACCGTT	CGTTCGCCC	AAGAAGGGGA	1560
TTTGGTTTTT	GCTAATGAAC	CGATTGTGCA	GGTGAAGGA	CCTCTAGCCC	AATGTCAGTT	1620
GGTCGAAACG	GCTCTTTTGA	ACATCGTCAA	CTACCAGACT	TTGGTGGCGA	CGAAGGCAGC	1680
TCGTATTCGT	TCGGTTATCG	AAGATGAACC	CTTGATGGAG	TTTGGGACAC	GTCCGGCTCA	1740
AGAAATGGAT	GCGGCCATCT	GGGGAACACG	CGCAGCTGTG	ATTGGTGGCG	CCAATGGAAC	1800
CAGCAACGTG	CGTGCGGGTA	AGCTCTTTGA	CATTCCTGTT	TTGGGAACCC	ATGCCCATGC	1860

CTTGGTACAG	GTTTATGGCA	ATGACTATGA	AGCTTTCAAG	GCTTACGCTG	CGACCCACAA	1920
AAATTGTGTC	TTTCTTGTGG	ATACCTATGA	CACCCTTCGC	ATCGGTGTAC	CAGCTGCCAT	1980
TCAGGTGGCG	CGTGAGCTGG	GTGATCAGAT	TAACTTTATG	GGTGTGCGGA	TTGACTCTGG	2040
GGATATTGCC	TACATTTCTA	AGAAAGTCCG	TCAGCAACTG	GATGAGGCTG	GATTTACAGA	2100
GGCTAAGATT	TATGCTTCTA	ATGATCTAGA	TGAAAATACC	ATCCTTAACC	TCAAGATGCA	2160
AAAGCCAAG	ATTGATGTCT	GGGGTGTGGG	TACCAAGCTG	ATTACAGCCT	ATGACCAGCC	2220
GGCTCTTGGG	GCGGTTTACA	AGATTGTTGC	AATCGAAGAT	GAAACTGGTC	AGATGCGCAA	2280
TACGATTAAG	CTGTCTAATA	ATGCTGAAAA	AGTTTCTACG	CCAGGTAAGA	AGCAGGTGTG	2340
GCGCATTACC	AGTCGTGAAA	AAGGCAAGTC	AGAAGGCGAC	TATATCACTT	ATGATGGTGT	2400
GGATATTAGC	GACATGACAG	AAATCAAGAT	GTTCCATCCG	ACCTATACAT	ACATCAAGAA	2460
GACGGTTCGT	AATTTTGATG	CCGTTCCTCT	CTTGGTGGAT	ATCTTCAAAG	AAGGAATATT	2520
AGTTTACAAC	TGCGCTAGTT	TGACTGACAT	TCAGGATTAT	GCCCCTAAGG	AATTTGACAA	2580
GTTGTGGGAT	GAGTATAAGC	GTGTGCTCAA	TCCGCAGCAC	TATCCAGTGG	ATTTGGCGCG	2640
TGATGTATGG	CAAGATAAGA	TGGACTTGAT	TGATAAGATG	CGCAAGGAAG	CCCTTGGTGA	2700
AGGAGAAGAA	GAATGAGTTT	GCAAGAAACG	ATTATCCAAG	AGCTGGGTGT	CAAACCAGTG	2760
ATTGATGCCC	AGGAAGAAAT	CCGTCGTTCT	ATTGATTTCT	TAAAAAGATA	TCTGAAAAAA	2820
CATCCCTTCC	TAAAAACCTT	TGACTAGGG	ATTTCTGGGG	GACAAGACTC	AACCTTGGCA	2880
GGACGTTTGG	CGCAATTAGC	TATGGAAGAA	CTGCGAGCTG	AAACGGGAGA	CGATAGCTAC	2940
AAATTTATCG	CTGTCCGCCT	GCCATACGGA	GTGCAAGCTG	ATGAAGCAGA	TGCTCAAAAA	3000
GCCCTAGCCT	TCATCCAGCC	AGATGTCAGC	TTGGTTGTGA	ATATCAAGGA	ATCAGCTGAT	3060
GCCATGACAG	CTGCAGTTGA	AGCGACAGGT	AGTCCGTGTT	CAGACTTCAA	CAAGGGGAAT	3120
ATCAAGGCAC	GTTGCCGTAT	GATTGCTCAG	TATGCCCTTG	CTGGTTCCCA	TAGCGGAGCG	3180
GTCATTGAA	CAGACCACGC	CGCGGAAAAT	ATCACAGGTT	TCTTTACCAA	GTTTGGTGAC	3240
GGCGGTGCGG	ATATTCTCCC	TCTTTACCGC	CTCAATAAAC	GCCAAGGAAA	ACAGCTCTTG	3300
CAGAAACTTG	GCGCAGAGCC	AGCCCTTTAT	GAAAAAATCC	CAACGGCAGA	CCTAGAAGAA	3360
GATAAACCAG	GCCTAGCTGA	CGAAGTCGCA	CTTGGAGTCA	CCTACGCAGA	GATTGACGAC	3420
TACCTAGAAG	GCAAAACAAT	CAGCCCAGAA	GCTCAAGCGA	CCATTGAAAA	CTGGTGGCAC	3480
AAAGCCAAC	ACAAACGCCA	CTTACCCATC	ACCGTATTTG	ATGACTTTTG	GGAGTAAAAA	3540
GGTCCGGGGG	ACCTTTTATG	CTTCTTGCCC	TGAAATTAAT	AAGCAAGAAA	AACCTCCACT	3600

918

GGAGGTTTTTC	AGCCTCTCAT	CTTGAAATAA	GAAAGTGAGA	GAAGGTCTGG	GGGATCTTGA	3660
ACCCCGAGTT	TAGAAATAAG	AAAATGAGGC	AGATTCAGTA	ACTCGAAGAG	TTCGATTTCA	3720
TCGTCTTACC	CCTGCAACGA	TGACTAGGTT	TGAAAAAGCT	TGCTAGAGCG	CATTTCAAAC	3780
CAGGCAGCAA	CTGCGTCAAG	AAATTAGAAG	ACAACTCGT	TTTCTAGCTG	TACTGAGTT	3840
GAGCCTTTTT	ACTACGAGTA	TAGAAATAAG	GAAGTGAGGT	AGCATCATGA	AATCTATCGG	3900
TACGCAAATA	TTACAGACAG	AACGTTTGAT	TTAAGAAGA	TTTGTGGAGA	GTGATGCAGA	3960
AGCCATGTTT	CAAAATTGGG	CTTCATCCGC	TGAGAATCTG	ACCTATGTTA	CCTGGGATCC	4020
CCATCCTGAT	GTCGAAATCA	CTCGAAACTC	GATTTGCAAT	TGGGTTGCTT	CCTATACTAA	4080
TCTCAACTAT	TATAAATGGG	CCATTGTCT	AAAAGAAAAC	CCAGAGCAAG	TAATAGGAGA	4140
TATCAGCATT	GTTAAGATAG	ACGAGGCTGA	TTAAGCTGT	GAAATGGCT	ATGTGTTAGG	4200
CAAGGCTTAC	TGGGGAAATG	GTATGATGAC	AGAGACTTTG	AAAGCTATCT	TGGACTTTTG	4260
TTTTACTCAA	GCAGGTTTTC	AAAAGGTCAG	AGCACGTTAT	GCCAGTCTCA	ACCCAGCTTC	4320
AGGTCGTGTC	ATGGAAAAGG	CTGGAATGTC	CTATCTACAA	ACCATTGTTA	ATGGGTGAGA	4380
GAGAAAAGGC	TATCTTGC	ATCTTATTTA	TTATGGTATA	AGTAGGGAAG	AATGTTGAAT	4440
TCTATTTTCT	TTTTCTATCG	AAGTCAACTA	TTTATGTAA	ATATAATAAT	TAGCATTTCCA	4500
AGTTTATTTG	AAACTTTAAA	ATAGCATATT	GATTAGTACA	AGACAGATGT	TCTAGTTCCT	4560
TCTTTAATCT	GGTTTAGTGT	TAGTTAAAAA	ATCGCTTTAA	GCTTGTA	AAGAGGGAGC	4620
TAATCGACTA	GATTCPCAG	CCGAACAGGI	GGTAATGTAC	TTTTTATAGT	GTAATCCTAG	4680
CTGTGTTAA	ATTTAAAATA	GAATCCTCTA	TCGAGTTAGG	GAATTA	CAACCAATTT	4740
TATTCATGTT	TTTTCTATCA	AATTATCTAA	TATTTAAAATA	GTCTCATTCT	GATGAGAAAA	4800
CTATTC	ATCATTCATA	CCTCTCTCAA	CTAGATGTAA	CTTACAAAAC	CCCTGACCTC	4860
ATGAGCCACT	TTCTTCTCC	TCATGAGGTC	AGTTTTACTT	TCTGCTGTTT	CAGTATCGTT	4920
TTTCTCGCT	AGATTTCC	AAAAGGGCAG	ACTCCTCCCT	TGGTGCGTCA	CACGATTTTT	4980
TCATCTCGAC	TGTTCTTTAA	TGCATCATT	ACGACGCTTT	TCTTCTAGGT	GGTTCATAAG	5040
GAACAGGAAG	ATTCAGGTTG	ACTTTTCTAA	TCCTAGAATA	AAGTGCTGAA	AACAATTCGG	5100
AATAGGCATA	GAGACTAGAC	AATTTGAGGA	GCTGCTTGC	TCCTGTTTCA	ACACATTTTC	5160
CCACCACGTG	AAGAAAAAGA	TGGCGGAAGC	GTTTGATTGT	TAAAGTTTGG	AAGTCACCTC	5220
CAGCTAGATG	TTTGAGAAAA	AGATAGAGAT	TGTAGGCGAT	ACAGCTCATC	ATCATACGAA	5280
CTTCGTTTTT	GATTAAGGTT	GA	ACT			5305

(2) INFORMATION FOR SEQ ID NO: 136:

919

- (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 3964 base pairs
 (B) TYPE: nucleic acid
 (C) STRANDEDNESS: double
 (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 136:

TGGCAGCTCG TCGTCGTAAA GGACGCAAAG TTTTGGCTGC ATAATCCAAA CGAATTCTAT	60
CAAAAATCAG TAGGAACTCG AGTCTACTGA TTTTATTFFF TGTA AAAAAG TTCAGTAGAT	120
GCAAATGGAT TCGGAAGCGA TGTTACAGTA GATTGAAACT AGAATAGTAC ACCTCTGTTT	180
CTAAAACATT GTTAGAAATC GATTTGACTG TCCTGATCGA TTTGTCCCTGT TATFATTTTA	240
TTTTACTATA AAGTTGAAGT AGGTGGAGAT GGTACAGCAA CAATCGTCTT TAAAGATGGT	300
TCAGCTATTA CAATTCCAGG AAATCAATTG GTAGCACAAAG ATCCAAAAGC ACAAGATAGC	360
ACTAAACTGA CTGCTGAAAA ATCAACTGTT AAAGCACCTG CTCAAAGAGT AGATGTAAAA	420
GATATAACTC ATTTAACAGA TGAAGAAAAA GTTAAGGTG CTATTTTACA AGCAAATGGT	480
TCAGCATTAG ACGGAGCGAC AATCAATGTA GCTGGAGATG GTACAGCAAC AATCACATTC	540
CCAGATGGTT CAGTAGTGAC GATTCTAGGA AAAGATACAG TTCAACAATC TCGGAAAGGT	600
GAATCTGTAA CTCAAGAAGC TACACCAGAG TATAAGCTAG AAAATACACC AGGTGGAGAT	660
AAGGGAGGCA AFACTGGAAG CTCAGATGCT AATGCGAATG AAGGCGGTGG TAGCCAGGCG	720
GGTGGATCAG CTCACACAGG TTCACAAAAC TCAGCTCAAT CACAAGCTTC TAAGCAATTA	780
GCTACTGAAA AAGAATCAGC TAAAAATGCC ATTGAAAAAG CAGCCAAGGA CAAGCAGGAT	840
GAAATCAAAG GCGCACCGCT TTCTGATAAA GAAAAAGCAG AACTTTTAGC AAGAGTGGAA	900
GCAGAAAAAC AAGCAGCTCT CAAAGAGATT GAAAAATGCGA AACTATGGA AGATGTGAAG	960
GAAGCAGAAA CGATTGGAGT GCAAGCCATT GCCATGGTTA CAGTTCCTAA GAGACCAGTG	1020
GCTCCTAATG CTGCTCCTAA GACAACAAGT GCACCGCAAG CAACTGCAGG AACAATGCAA	1080
GATGTTACCT ACCAGTCACC TGCTGGCAA CAATTACCTA ACACAGGTC AGCATCAAGT	1140
GCAGCACTG CTAGTCTTGG TCTAGTGGTG GCAACAAGTG GTTTGCTTT GCTAGGAAGA	1200
AAGACTAGAC GTAGAAAATA GAACAGCTAG AAAATTCTAT TCTCTACTTA AAGTTAGATT	1260
ATAAGGGGGA TTTTGAGAAG TCATCAATCC TAGTGATGGG TGAGAAAAGT GAGAACCCAA	1320
GATAATCACA TACTTTAGCT GAATAGGAAT ATTCTATCAA TGTAGCCAAT CTCTTCTGTC	1380
TCTAACTGTG GAATAGGAGA TGGGCAATAT CGGATAGAAA AGATAGCAGA ATAGCTCTCT	1440

ATTGAAGAGA	GGAGGGGAAA	CCGAAAAATT	AGGTGCCCCCT	CCTCTTTTTT	GGTATAATAG	1500
AAGATAGAAA	ACGAGGTTAG	AAGAGATGAT	TTTTGATACA	CATACACACT	TGAATGTAGA	1560
AGAATTTGCA	GGTCGTGAGG	CAGAAGAAAT	TGCCTTGGCT	GCTGAGATGG	GTGTGACACA	1620
GATGAATATT	GTTGGTTTTG	ATAAACCGAC	GATTGAGCAT	GCCTTGGAGT	TGGTAGATGA	1680
GTATGAGCAG	CTCTATGCGA	CTATTGGTTG	GCATCCTACA	GAAGCTGGTA	CTTATACAGA	1740
GGAAGTTGAG	GCTTACTTGT	TGGATAAGTT	AAAACATTCC	AAGGTTGTGG	CTTTAGGTGA	1800
AATTTGGCTTA	GATTACCATT	GGATGACAGC	GCCCAAAGAG	GTGCAGGAGC	AGGTTTTTCG	1860
CCGTCAGATT	CAGCTATCTA	AGGACTTGGA	TTTGCCTTTT	GTTGTCCATA	CCCGTGATGC	1920
GCTGGAAGAT	ACCTATGAGA	TTATCAAGAG	TGAGGGCGTT	GGTCTCGTG	GTGGTATCAT	1980
GCATTCATTT	TCAGGGACGC	TTGAGTGGGC	AGAGAAGTTT	GTGGATCTTG	GTATGACCAT	2040
TTCCTTCTCA	GGAGTGGTGA	CTTTAAGAA	GGCAACTGAC	CTCCAAGAAG	CAGCTAAAGA	2100
GTTACCTTTG	GACAAGATGT	TGGTGGAAC	AGATGCGCCT	TACTTAGCAC	CTGTACCCAA	2160
GCGTGGTCGT	GAAAATAAAA	CAGCCTATAC	TCGCTATGTG	GTCGACTTTA	TCGCTGACTT	2220
GCGTGGTATG	ACGACAGAAG	AGCTGGCGGT	AGCAACGACT	GCAAATGCAG	AACGAATTTT	2280
TGGACTGGAC	AGCAAGTAAT	GAAAGAGAAA	ATTTCTCAAG	TTATCGTGGT	TGAAGGGCGT	2340
GATGATACGG	TCAATCTCAA	ACGTTATTTT	GATGTGGAGA	CCTATGAGAC	TCGAGGTTCT	2400
GCCATCAATG	CTCAGGATAT	AGAGCGGATT	CAGCGCCTGC	ACCAACGTCA	TGGAGTCATT	2460
GTCCTTTACAG	ACCCAGATTT	TAATGGGGAA	CGGATTCGGC	GCATGATCAT	GATGGTCATT	2520
CCAACAGTTC	AGCATGCCTT	TCTCAAGCGA	GATGAAGCTG	TTCCCAAGTC	CAAGACCAAG	2580
GGGCGTTCTC	TGGGAATTGA	GCATGCCAGC	TATGAAGACC	TGAAAACGGC	TCTAGCTCAA	2640
GTGACAGAAC	AATTTGAACA	TGAGAGTCAG	TTTGACATTA	GTCGTAGCGA	TTTGATTTCG	2700
CTTGGTTTTT	TAGCAGGGGC	AGACAGCCGT	AAGCGTAGAG	AATATCTCGG	AGAGACTCTC	2760
CGAATCGGCT	ATTCCAACGG	CAAGCAACTC	CTCAAACGCC	TAGAGTTGTT	TGGGGTACT	2820
TTGGCAGAAG	TGGAAGAAGC	TATGAAATCT	TATGAGTAGG	AAAGATGTAG	CCGTTACAAT	2880
TTTTTAAGTT	TCACAGTATT	TTTCGAAGCA	GGTAGAAGAG	GAGGCGTCTG	ATGTTAATTG	2940
GTCAAAAAAT	TAAAGAGATT	CGGATAGAAA	AAGGAATTAG	TCGTCCAGAT	TTTTGTGGAG	3000
ATGAGCAAGA	ACTGACAGTT	CGTCAACTGT	CGCGAATTGA	AAGTGGAGCT	TCGCAACCGA	3060
GTTTGCCCAA	GTTAGACTAT	ATTGCTCGCC	GGCTAGGAGT	TCCAGTTTAT	AGCCTTATGC	3120
CGGATTTTTT	AGCTCTTCCT	TCTGCTTATT	TAGAATTGAA	ATACCAGATT	TTACGTGAAC	3180
CAATCTATGG	TAAAGAAGAG	GAGTACGATA	AGAAGGAAGC	GTGTTTGAA	GAGATTTATA	3240

921

AAACATACTT	TGATAATCTT	CCTAAAGAAG	AACAATTAGC	ATGTGAAGTA	TTGCAGGCGT	3300
GTTTGGATAC	TTCTAGAACT	AGAAGCCTG	AATATGCAGA	GTTAATACTT	GAGGAACATA	3360
TGCCTCAGAT	TATAGAAAAA	GAAGCTTATT	CAATAAATGA	TATGTTGTTG	ATTTCGTTTGT	3420
TTTTTTATCA	AATGCTCATT	AGAAAAGATC	TTGCCAAAT	TATAAATCAA	ATCGAAAAGC	3480
TAATGCTCTT	TCTTTTGGA	CAGAAGAAGG	TAACTCAAAT	AGAGAATTAC	TTTATAATTA	3540
GAGATACTCT	TATTCAGGA	ATGTGTTGTC	TTGAAAAGGT	AGGAGTAACT	GATTGTTTTA	3600
ATGATTATCT	ATCGTGTTTA	CAAGAAATTA	TGGATAAAAC	TCAAGATTAT	CAAAGAAGAC	3660
CTCTTGATAT	TATGTTTTTG	TGGAAGCAAG	CATTAAGAGA	AGAAAGAGAT	TTTAGTTTAG	3720
CTGAATCAT	TTATCAGTCT	TCTAAAACAT	TTGCCGAGCT	AATTGGAGAT	GAATTTCTAG	3780
TAAAGAAAT	GACAGAGGAA	TGGCAAGAGG	ATGTCAAAAA	ATATTTATAA	ACATAGTGAA	3840
TCAGTGACAA	AGATGTCCTT	GTCCTCGTAT	CAAAACAGTT	CTAAAGTTCG	TCTTTAGGGA	3900
TGTTTTTTTA	GATATAAGCT	AAAAATGACA	CGAAATGTTT	AGATTTTAAG	GACATGATG	3960
TCCG						3964

(2) INFORMATION FOR SEQ ID NO: 137:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 12666 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 137:

TGAGACCGT	ATTTGTATTA	GGGAAATGGG	TATCTATTTT	TAATGCTGTG	GGGATTTTGA	60
TTGTTTCTAT	TATTCAAACC	AAAAGCTTGT	CAGGTATTGG	AGCAGGATTG	TTTAATCTAT	120
ATAACATTC	ATCTTATATA	GGTGATTTAG	TTAGTTTCAC	TCGATTGATG	GCATTAGGAT	180
TATCTGGAGC	AAGTATAGCA	TCAGCTTTCA	ATTTAATTGT	TGGTTTGT	CCGGGAATAT	240
TGGCTAAACT	GACAATTGGA	TTAGTATTAT	TCATCTTTT	ACATGCGATC	AATATTTTTT	300
TATCGTTACT	ATCAGGATAT	GTTCATGGAG	CACGCTGAT	ATTTGTTGAA	TTTTTTGGTA	360
AGTTTTATGA	GGGTGGAGGA	AAACCATTC	AACCTTGAA	GGCTTCTGAG	AAATATATTA	420
AGGTTATTAC	AAAGAATTAA	TGGAGGATAT	ATATAATGGA	ACATTTAGCA	ACTTATTTTT	480
CAACCTATGG	AGGAGCTTTC	TTGCTGCAT	TGGGAATTGT	ATTGGCGGTT	GGATTAAGCG	540
GTATGGGGTC	TGCTTATGGA	GTTGGTAAGG	CTGGGCAATC	TGCCGAGCT	TTACTGAAAG	600

AACAGCCTGA	AAAGTTTGCC	TCAGCTTTGA	TATTGCAATT	ATTGCCCGGA	ACACAAGGAT	660
TATATGGTTT	TGTATTGGA	ATTTAATTT	GGTTGCAATT	AACTCCAGAA	CTTCCTTTAG	720
AAAAAGGCGT	TGCTTATTT	TTTGTAGCTC	TTCCAATTGC	TATTGTAGGA	TACTTTTCAG	780
CTAAGCATCA	AGGAAATGTA	GCAGTAGCGG	GAATGCAAAT	CTTGGCTAAA	AGACCAAAAG	840
AATTCATGAA	GGGAGCAATT	TTAGCTGCCA	TGGTAGAAAC	CTATGCAATT	CTTGCTTTTG	900
TCGTATCATT	CATTTTGACC	CTTCGTGTAT	AAGAAATAAA	TTTGCAATTC	AAAGGAGGTG	960
TCTAAATGAG	CAATTTAGAA	AACTTACGAG	AGTCTGTTAT	TGAACAAGCT	CATGAAAAAG	1020
GGCGTATGAA	ATTATTGGAT	TCCAAAAAGA	AGATTGATGA	TGAATTTGAA	ATGCAAAAAGT	1080
CGCTCATTAT	AAAGAAAAAA	GAAGCTGAAC	ATGAACGAAA	GTTAAAAGAA	TTGCAACAGA	1140
AATATCAAAT	AATTTTCAA	CAATTAATAA	ATAAGGAACG	CCAATCAACG	TTAGTATCAA	1200
AACAGAAAAT	ATAAAAGAA	CTTTTCAAT	CTGCTTTACT	AGAAATGGAA	TCTTGGAGTG	1260
CAGATAAAGA	AATGGAGTTC	ATCTATCGAA	TTCTGGAACG	ATATTCACAA	CAAGAGGTCA	1320
TAGTAACCTT	TGGGGAACGG	ACTTTAGCTA	AATCAATTT	GGAACAATTA	GAGAAATTGA	1380
AATTCTCTTT	TCCAATTAT	TTATTTAGTG	AACAACCTAT	CTCAAATGAA	TCAGGCTTAC	1440
TTATTTCAAT	AGGTAAAAT	GATGATAACT	ATTTGTATAA	AACATTAATT	GGATCGATTT	1500
CTAAGGAAGA	AAGTTCAAGT	ATCGCAAATC	AAATTTTAT	CAATTAAGGA	TGAAATTGGT	1560
TAATCCTTCT	TAGAAATTTG	GAGTATTCCA	ATAAAATTAG	AAAGGTATTT	TATGGATACT	1620
AATCTTTTTT	CAAAAATAAA	TACGACGATT	TCGGTAAAAG	AAAACGATTT	TATTACAGAA	1680
GAAAAATTC	AAAAATTAT	ACAATCCAAA	GATACGGAGA	CATTGGCATT	TATCTTAGAA	1740
TCAACTCCCT	ATCATTTATC	GATTGACATC	TTAGAAGATC	CTAGTCAGAC	AGAGATTTCG	1800
CTAATGACAA	AATTAGTCAA	TGATTATAGA	TGGGCCTATG	CTGAAAGTCC	GTCTGATATA	1860
ATTGTGACTT	TATTTGCTTT	ACGATATGTT	TATCATAATA	TCAAAGTTTT	ATTAAAATCT	1920
AAGGCGGCAA	TTAAGAAAGA	TTTTTCTAAA	TTATTAATTC	CAATAGGGAT	TTTTGATATA	1980
GAAAGTTTAA	AACATTTAGT	TTCTTCCTTA	CATTCAGATA	CACTTCCTGA	TTTTATGGTT	2040
CGTGAAGTAG	AATCAATTTG	GAATGAGTAT	GAACTTTTA	ATAATATTCG	TGFACTTGAT	2100
GTCGGAGCTG	ATCTAGCATA	TTTAAACAT	CTGAACTTT	TATCTAATGA	GTTAGATGAG	2160
GTAAGTCTC	AGTTTATGTT	CGAAATGATT	GACTTTTATA	ATATTATTAC	TGTAACACGT	2220
GGTTTATCTC	AAAATAAGAG	TCATGGGGAT	ATTTTACAAT	TACTTTTCAGA	TGAAGGAAGT	2280
ATTTCTGCTA	AAGAATTTAT	ATACATTGTA	GAAAATCAAG	AAATATTTGT	GTGGTTCAAT	2340
AAAATAAATC	CAAGCTTAGA	TTCAATCTTT	TCAACTTATG	AATTGAAGAT	GCAGGACGCA	2400

ACAATTCAT	CTTCTGAGTT	AGAATTTT	TGTGATTTAC	TATTGTATAA	AACTTTAGAT	2460
CAAGGAAGGT	ACAATGTAGA	GGGGCCGTTA	GTTCTTGCTA	GATATTTATT	GGGATGTGAG	2520
TTTGAAGTAA	AGAATCTCAG	AATGATCATA	TCAGCTCTTC	AAAATACAAT	TCCCTTTGAA	2580
TCAATAAAAG	AAAGGATACG	CCCACATTAT	GGAAGCTAAT	AAGTATAAAA	TTGGCATAAT	2640
TGGTAGCCGT	GATATTATTT	TACCATTTAG	CATGATTGGG	TTTGATATAT	TTCTGCCTA	2700
CCAAGAACAA	GAAGCTATAA	ATACACTAAG	AAAATTAGCT	CAATCTGATT	ATGGTGTCAT	2760
TTATATCACT	GAAGACATTG	CTTCAATGAT	ATTAGATACA	ATTCGCCATT	ATGATTCCCA	2820
AGTTGTGCCT	GCTATTATTT	TATTACCGAC	TCATAAACAA	GGTTTAAATT	TAGGATTAAG	2880
ACGTATAGAG	GATAATGTAG	AGAAAGCAGT	AGGACACAAT	ATTTTATAAT	AATGTACAAA	2940
ATTGTCTGTA	ATATTATTCT	ATAATTTT	GACTTAGTAA	GGAGAATAAC	TTTGACTCAA	3000
GGGAAGATTA	TAAAAGTATC	GGGACCTCTA	GTTATTGCAT	CAGGTATGCA	GGAGGCTAAT	3060
ATTCAAGATA	TTTGCCGTGT	AGGTAAGCTA	GGGTTAATCG	GTGAAATTAT	TGAAATGAGA	3120
AGAGATCAGG	CATCTATCCA	AGTCTATGAA	GAAACATCTG	GTCTTGGTCC	GGGAGAACCT	3180
GTTGTTACAA	CTGGAGAACC	TCTCTCGGTT	GAATTAGGGC	CAGGATTGAT	TTCTCAAATG	3240
TTTGATGGCA	TACAACGCCC	ATTAGATCGA	TTTAAATTGG	CTACTCATAA	TGATTTTCTA	3300
GTTTCGTGGG	TAGAAGTTCC	AAGTTTGGAT	AGAGATATTA	AGTGGCATT	TGATTCCACT	3360
ATAGCAATTG	GTCAAAAAGT	GAGTACGGGT	GATATTCCTG	GAACGTCAA	GGAAACCGAG	3420
GTAGTAATC	ATAAAATTAT	GGTTCCTTAT	GGAGTATCTG	GAGAAGTCGT	TTCTATTGCA	3480
TCTGGCGATT	TTACAATTGA	TGAAGTTGTA	TATGAAATAA	AAAAATTGGA	CGGTAGTTTC	3540
TATAAAGGAA	CGCTTATGCA	AAAATGGCCT	GTCCGCAAGG	CGCGTCTGT	TTCTAAACGT	3600
TTAATTCCAG	AAGAACCATT	AATCACAGGT	CAACGAGTTA	TTGATGCATT	CTTTCCAGTA	3660
ACCAAAGGGG	GAGCTGCAGC	AGTTCCTGGA	CCGTTTGGAG	CAGGAAAGAC	AGTTGTACAA	3720
CACCAAGTAG	CTAAATTTGC	CAATGTTGAT	ATTGTTATTT	ATGTCGGTTG	TGGAGAACGT	3780
GGAAATGAAA	TGACGGATGT	ACTGAATGAG	TTTCTTGAGT	TGATTGACCC	TAATACCGGA	3840
CAATCAATTA	TGCAACGGAC	AGTTCCTGAT	GCTAATACTT	CAAATATGCC	TGTTGCTGCT	3900
CGTGAGGCTT	CAATTTATAC	AGGAATTACC	ATGGCTGAGT	ATTTTCGTGA	TATGGGCTAC	3960
TCTGTCGCCA	TTATGGCTGA	TTCAACTTCA	CGTTGGGCAG	AAGCGCTACG	TGAAATGTCA	4020
GGACGTCTAG	AAGAAATGCC	TGGTGATGAG	GGTTATCCTG	CTTATCTGGG	AAGTCGTATC	4080
GCTGAATATT	ATGAAAGAGC	AGGACGTTCT	CAGGTTCTAG	GGCTTCCAGA	ACGTGAAGGA	4140

ACGATTACTG CTATTGGAGC TGTATCGCCA CCTGGTGGAG ATATTTTCAGA ACCAGTTACT 4200
 CAAAACACTT TACGGATTGT GAAAGTTTTT TGGGGGCTTG ATGCTCCGTT GGCACAGCGA 4260
 CGTCATTTTC CTGCAATTAA CTGGCTTACA TCTTATTCAC TATATAAAGA CAGTGTGGGC 4320
 ACTTATATAG ATGGTAAAGA GAAGACAGAT TGGAAATAGTA AAATAACTCG TGCGATGAAC 4380
 TACTTACAAC GGAATCTAG TTTAGAGGAA ATTGTTTCGTC TTGTTGGAAT TGATTCTCTG 4440
 TCTGATAATG AACGACTAAC GATGAAAATT GCTAAACAAA TTCGAGAAGA TTATTTGCAA 4500
 CAGAACGCTT TTGATTCCGT AGATACATTC ACTTCGTTTG CAAAACAAGA AGCAATGCTA 4560
 AGTAATATTC TCACTTTTGC TGATCAGGCA AATCATGCTT TAGAGTTGGG TTCTTACTTT 4620
 ACAGAGATTA TGAAGGTAC CGTGGCAGTT CGAGACCGTA TGGCGAGAAG TAAATATGTT 4680
 TCAGAAGATA GATTAGATGA AATCAAAATT ATATCAAATG AGATTACACA TCAAATTCAT 4740
 TTGATATTAG AAACAGGAGG TCTATAAATG AGTGTATATA AAGAATACAG AACTGCTAGT 4800
 GAAGTTGTTG GGCCTCTTAT GATTGTTGAA CAAGTAAATA ATGTGTCTTA CAATGAGTTA 4860
 GTTGAAATTC AACTTCATAA TGGAGAAATT CGTCGTGGAC AAGTTTTAGA GATCCACGAA 4920
 GATAAAGCAA TGGTTCAGCT TTTTGAAGGA TCTAGTGGAA TAAATTTAGA AAAGTCTAAA 4980
 ATTCGTTTTG CTGGTCATGC ATTAGAATTG GCTGTATCTG AGGATATGGT TGGTCGTATT 5040
 TTTAATGGGA TGGGAAAACC AATTGATGGT GGACCAGATT TAATTCCAGA GAAATATTTA 5100
 GATATTGATG GTCAAGCTAT TAATCCTGTA TCTAGAGATT ATCCAGATGA ATTTATTCAG 5160
 ACAGGGATCT CCTCTATPGA TCATTTGAAT ACTCTTGAC GTGGTCAAAA ATTACCAGTA 5220
 TTTTCAGGTT CGGGCTTACC TCATAATGAA TTAGCTGCTC AGATAGCAAG ACAAGCGACT 5280
 GTTTTAAATT CTGATGAAAA TTTTGCGGTT GTATTTGCAG CAATGGGTAT TACTTTTGAA 5340
 GAAGCTGAGT TTTTATGGA AGAACTCAGA AAAACAGGAG CGATCGATCG TTCGGTTTTA 5400
 TTTATGAACT TGGCAAATGA TCCTGCAATT GAGCGTATTG CAACTCCCCG CATTCCTTTA 5460
 ACTGCGGCAG AGTATCTAGC TTTTGAAAAA GATATGCACG TTCTAGTTAT CATGACGGAT 5520
 ATGACTAACT ATTGTGAAGC GTTACGTGAA GTCTCGGCAG CTCGCCGTGA AGTTCAGGG 5580
 AGACGAGGCT ATCCGGGATA TTTATATACA AATTTATCAA CTCTATACGA AAGGGCTGGT 5640
 CGCTTAGTTG GTAAAAAAGG TTCGGTGACA CAGATTCCTA TTTTAACAAT GCCAGAAGAT 5700
 GACATAACAC ATCCAATTCC TGATTTAACT GGATACATTA CTGAAGGCA AATTATTTTG 5760
 TCGCATGAGT TGTATAATCA AGGTTATCGT CCACCAATCA ATGTTTTACC TTCTCTCTCT 5820
 CGATTAAAAG ATAAGGGATC TGGAGAAGGT AAAACTCGTG GAGATCATGC TCCAACATG 5880
 AATCAACTGT TTGCAGCCTA TGCCCAAGGG AAAAAGGTTG AAGAGTTAGC AGTAGTATTA 5940

925

GGAGAATCGG	CTTTATCTGA	TGTAGATAAA	TTGTATGTGA	GGTTTACAAA	GCGTTTTGAA	6000
GAAGAGTACA	TAAACCAAGG	ATTTTATAAA	AATCGAAATA	TAGAAGATAC	GTTGAATCCT	6060
GGGTGGGAAT	TACTATCAAT	TCTTCCTAGA	ACAGAGTTAA	AACGTATCAA	AGATGATTTG	6120
CTTGATAAAT	ACTTACCTTT	GGTAGAAGTT	TAATCCGAA	ATGGAGTGAT	TATCTATGGT	6180
ACGTTTGAAT	GTAAAACCAA	CTCGTATGGA	ATTGAATAAC	TTAAAGGAAC	GTTTGACAAC	6240
AGCTGAACGT	GGACATAAGT	TATTAAAGGA	TAAAAGAGAT	GAATTGATGA	GGCGATTTAT	6300
TTCTTTGATT	CGTGAGAATA	ATCAACTTCG	GAAAGAAGTG	GAAAGTTATC	TAATTGATAA	6360
TCTAAAATCC	TTTGCAGTTG	CTAAATCATT	AAAGAATTCT	CAAATGGTGG	AGGAATTATT	6420
TTCAATTCCA	TCGAAAGAAA	TTGAATTATT	TGTTGAGAAA	GAAAATATCA	TGAGTGTAAC	6480
AGTTCCTAGA	ATGCATATGA	ATATTACTTC	TCAAAATGAG	AACAGTGAAT	ACAGCTATTT	6540
ATCTTCTAAT	AGTGAAATGG	ATGATGTATT	TGCTACAATG	AATAGTTTAA	TTTATAAATT	6600
ACTAAGACTG	GCAGAAGTTG	AAAAACGTG	TCAGTTAATG	GCTGATGAAA	TAGAAAAAAC	6660
ACGTAGACGT	GTAAATGGTT	TAGAATACTC	GATTATTCCA	AACCTGTCGG	AAACTATFCA	6720
TTATATAGAA	TTGAAACTAG	AGGAGGCAGA	AAGAGCCAAT	TTAGTTCGTA	TTATGAAAGT	6780
GAAGTAGATC	CTTTATTTAG	ATTATTAATT	AGATGAACAA	ATATCAGCTT	GGATAAGGCT	6840
TTAAGCCTTT	CTAAGCTTTT	TTTATTGACA	GTATCAGGAT	ATCTTTTCA	AAATTTTGGT	6900
TTGTTAGATA	ATGAAAATGT	TTCTACTAAT	CTAGATTTAG	GATTAGTAAA	TCGTAAATGT	6960
AATTATATAG	AAAGTAAGCG	CGTCATAACA	AGGTATCTAT	CATTCATGGA	GCTCCTCCTG	7020
TATACTATTA	GTAAAGTAAA	ACTATTGGAG	GATATTTTAA	TGCCACAACC	TATTGTTCCCT	7080
GTAGAGATTC	CACAATCTCG	TCGTTTTGAT	TCTAAAAGA	GAAATGATAT	TCTGCTTAAA	7140
ATTTCGTATTG	GCAAGCTTGA	AGTAAGTTTT	TTTCAATCTC	TCAATCTCGA	AATGGTAGAA	7200
CAGCTTTTGG	ATAAGTGTT	GCTCTATGAC	AATTCATCTA	TCTAGCCTAG	GGGAGGTCTA	7260
TCTCGTGTGT	GGGAAAACCTG	ATATGAGACA	AGGAATCGAT	TCACTGGCTT	ATCTGGTTAA	7320
AACCCACTTT	GAATTGGATC	CTTCTCCGG	TCAAGTCTTT	CTCTTTTGTG	GTGGACGTAA	7380
AGACCCTTT	AAAGTCCTTT	ACTGGGATGG	TCAAGGATTT	TGGCTACTAT	ATAAACGCTT	7440
TGAGAACGGC	AGATTGATTT	GGCTAAGTAC	AGAAAAGGAT	GTCAAAGCTC	TCACACCAGA	7500
ACAAGTAGAC	TGGCTTATGA	AGGGCTTTTC	TATCACTCCA	AAAATATAGT	AGATTGAAAC	7560
TAGAATAGTA	CACCTCTGCT	TCTAAAACAT	TGTTAGAAAT	CGATTTTACT	GTCCTGATCG	7620
ATTTGTCCTG	TTCTTATTTT	ATTTTACTAT	AAATCCATCA	GAAAGTCGTG	ATTTCTATTG	7680

AAATGAGGAC	TTTCTTTT	TACTCATCTG	CTTTCAAAAA	GCATTCTAGT	CCATCTCCGA	7740
TTAACGATGG	ACTTTATCAC	CTCCTTCTCC	AGTCCTTGTA	TAACATCTTG	GAGTTGATTC	7800
ATGACATCTT	CCAAAGTTTA	AAAGGCTTTA	TTCTTAAATC	CACGTTTACG	AATCTCTTTC	7860
CACACTTGTT	CAATGGGGTT	CATCTCTGGT	GTGTATGGAG	GAATAAATGC	AAAGCCAATA	7920
TTAGTCGGAA	TCTTTAAGGT	ACTTGATTTA	TGCCATATAG	CATTGTCCAT	AACGAGTAAA	7980
AGATAATCAT	CTGGATAAGC	TTGTGAAATC	TCCTATTCCT	AAAGCCCCTT	TAGCGCATAA	8040
CTTTGGCTCA	GCTTCTATTA	TCGCTCACAC	CATCCATCAG	AAGTTTAATC	TGAAGGTACC	8100
CAATTATCGC	CAAGAAGAAG	ATTGGGCTAG	GATGGGTTTA	CCAATCACAC	GTAAGGAAAT	8160
CTCTAATTGG	CATATCAAGG	CGAGTCAATA	CTATTTGGAG	CCCCTTTATA	ACCTCTTGCG	8220
AGAGAGACTA	TTGACTCAGC	CCTTACTTCA	TGCGGATGAA	ACTTCTTATA	GGGTGCTAGA	8280
GAGTGATAGT	CAGCTGACTT	ACTATTTGGAC	TTTTTTGTCA	GGTAAAGCAG	AGAAACAAGG	8340
GATTACGCTT	TACCACCATG	ATCAGTGTCTG	AAGTGGTTCA	GTAGTACAAG	AATTCCTAGG	8400
AGATTATTCT	GGCTATGTGC	ATTGTGATAT	TTTGCGGCAG	TAACTTAGGA	CTTTAGTCCT	8460
CTAGTTCTGC	CTATGCGATA	GCAGTCCAAG	GTTTAGGAGC	AAGGCGACGC	TAAGCTTGGT	8520
AAACTTCGAA	CCGCTCGTCT	GCTTATCGTC	AACTGGAAGA	AGCTGAACTT	GTTGGATGTT	8580
GGGCGCATGT	GAGAAGGAAG	TTTTTTGAAG	CGCCCCCCA	AGCAAGCGGA	TAAATCATCC	8640
TTAGGAGCTA	AAGGTTTAGC	TTATTGTGAT	CAGTTATTTT	CCTTGAAAG	AGACTGGGAG	8700
GCTTTGCCAG	CTGATGAACG	ACTACAGAAA	CGTCAAGAAC	ATCTCCAGCC	CTTAATGGAA	8760
GACTTCTTTG	CTTAGTGCCG	GCGTCAGTCA	GTTTTAGCAG	GTTCAAACT	AGGAAGGGCA	8820
ATTGAATACA	GCCTCAAGTA	TGAAGAAACC	TTTAAGACCA	TTTTGAAAGA	CGGACATCTG	8880
GTCCTTTCCA	ATAATCTAGC	TGAACGCGCC	ATTAAATCAT	TGTTATGGG	ACGGAGTAAA	8940
AGAGTCCAGT	GGACTCTTTT	AGCCTAAGCT	CAGTTTAAAA	AAGCGAGGGT	GGTTATTTTC	9000
TCAAAGTTTT	GAAGGAGCTA	AAGCAAGAGC	TATTATTATG	AGTTTGTGG	AAACAGCTAA	9060
ACGTCATCAA	TTAAATAGCG	AGAAATATCT	ATCCTATCTT	CTAGAATGTC	TTCCAAACGA	9120
GGAAACTCTC	GTAACAAAG	AGGTTTTAGA	GGCTATTTA	CCATGGACTA	AAGTTGTACA	9180
AGAAAAGTGC	AAATAAGAAA	TCTCCAGATT	AGGAACTATC	CGTGAGTTCT	CCAGTCTGGA	9240
GATTTTTCAA	TAGACTTCCT	GCGAAACAAA	ATATGGTATA	ATAGTTCTAT	GAATGATGAA	9300
GCAAGTAAAC	AACTAACCGA	TGCACGATTT	AAGCGTCTTG	TTGGTGTTC	ACGCACGACT	9360
TTTGAAGAGA	TGTTAGCTGT	ATTAAAAACA	GCTTATCAAC	TTAAACACGC	AAAAGGTGGA	9420
CGAAAACCTA	AATTAAGTCT	AGAAGACCTT	CTTATGGCCA	CTCTTCAATA	TGTGCGAGAA	9480

TATCGAACTT	ATGAACAAAT	TGCGGCTGTT	TTTGGTATTC	ACGAAAGCAA	CTTAATCCGT	9540
CGGAGCCAAT	GGGTTGAAGT	AACTCTTGTT	CAAAGTGGTG	TTACGATTTT	AAGAACTCCT	9600
CTCAGTTCG	AGGACACGGT	AATGATTGAT	GCGACGGAAG	TAAAAATCAA	TCGCCCTAAA	9660
AAAAGAATTA	GCGAATTATT	CTGGTAAAAA	GAAATTTTAC	GCTATGAAGG	CTCAAGCGAT	9720
TGTCACAAGT	CAAGGGAGAA	TTGTTTCTTT	GGATATCACT	GTGAACATTT	GTCATGATAT	9780
GAAGTTGTTC	AAAATGAGTC	GCAGAAATAT	CAGACAAGCT	GGTAAAATCT	TGGCTGACAG	9840
TGGTTATCAA	GGGCTCATGA	AGATATATCC	TCAAGCACAA	ACTTCACGTA	AATCCAGCAA	9900
ACTCAAACCG	CTAACAATTG	AAGATAAAGT	CTATAACCAT	GCGCTATCTA	AGGAGAGAAG	9960
CAAGGTTGAG	AACATCTTTG	CCAAAGTAAA	AACGTTTAAA	ATGATTTTCAA	CAACCTATCG	10020
AAATCATCTA	AACGCTTCGG	ATTACGAATG	AATTTGATTG	CTGGTATTAT	CAATCATGAA	10080
CTAGGATTTCT	AGTTTTGCAG	GAAGTCTATT	ATCAAAAATA	CCATCAAGAT	TATATAAGAT	10140
TGATACAGGA	AAAGTTTTAT	TTGATGGTGT	AAATATTAAT	CAAATAGATA	AAAAAATATT	10200
AAGTCAAAT	TTAGGAGTAG	TTCCACAGGA	TTCAATTTTA	TTGAACCGAA	GTATTTCTTGA	10260
TAATATAACT	TTAAAGCACG	AAGTTACTTC	ACAAAAGATA	GAGGAAGTTT	GTAAAGCAGT	10320
TCAAATCTAT	GATGAAATCA	TGGCTATGCC	GATGAAATTT	AATACTATCA	TCTCAGAGAT	10380
GGGGTCAAAT	ATTTACAGTG	GGCAAAGGCA	ACGGATAGCA	CTGGCACGTG	CATTAATAAA	10440
TAATCCTAGT	ATTGTAATTT	TAGATGAAGC	AACTAGTGCA	TTAGACACTA	TTAATGAGGA	10500
AAGAATAACA	AAGTATATAC	AAAGTCAGGG	CTGTACTCAA	ATAATGTAG	CTCATAGATT	10560
GTCAACGATT	AAGGATGCGG	ATGTTATTTT	TGTAATGAAA	GGTGGTAAGA	TTGTTGAGTC	10620
AGGAAATCAT	AAGTACTTAA	TGGATCTTGG	TGGAGAGTAC	TACAGCTTAT	ATACAAAAG	10680
GAAATGAGGT	GTAAGAAAA	TGAAGAAAGA	AAATGAATAT	GTAATTTTAA	CAACAGCCTC	10740
ACTAGGGGTG	ATGATTGGAA	TAGTGTTTGC	AATTTTTTTA	GATTTTCCAG	TTGAATATGG	10800
TATTTCTTTA	GGCTTGTTGA	ATGGAATAGT	ATTGGGTTCG	CTGATTGTTT	ACAAAACAA	10860
TAAGAATTA	GCATAATTTT	TTGCTGTAAA	CTAAGGAGTA	GAGATGGCTA	TAGTTGAAAT	10920
TATAAATCTA	ACAAAAGCT	TTAAAGATAT	TGAAGTTATT	CATAACACTT	AAATAATAGA	10980
GCAACTACAG	TAGTAGCTTA	AAAACATGAT	TAAATCGCTA	TTCTTAGGAG	TAGCGGTTTT	11040
TCTTTTGTG	TAATACTCTT	TGAAAATCTC	TTCAAACCAC	GTCAGCTTTG	CTTTACCGTA	11100
CTCAAGTACA	GCCTGCGGCT	CGCTTCCTAG	TTTGCTCTTT	GATTTTCATT	GAGTATAAAA	11160
AGGGTCAAGT	AAGTATAGTA	AATTGAAATA	AGATATGAAC	AAATCGATTA	GAAAAGTCAA	11220

928

ATFAATTCT	AGAAATATGT	TAGAAATTGG	TTTGAATTCC	GCAATCAATT	TGTCAGTTT	11280
TTATTTTCATT	TCATTTTATT	TAATTAGATT	TTCCAATTTT	TTAATTCAAG	CTAAAAATCC	11340
CCAATCGTAG	TGATTGAGGA	TTGAGTAAAT	AAATCTTAAA	CAATACCTTG	TGCAATCATG	11400
GCATTTGCTA	CATTTTCAA	GGCAGCAATG	TTAGCTCCTG	CAAGGTAGTC	TTTATCAAGA	11460
CCGTATGTTT	CTGAAGTCGT	TTTAGCTGTG	TTGAAGATGT	TTGTCATGAT	GTCFTTGAGA	11520
CGGCCATCAA	CTTCTTCACG	AGTCCATGAG	AGGCGAAGAC	TGTTTTGGCT	CATTTCAAGA	11580
GCTGAAACGG	CTACACCACC	AGCGTTGGCA	GCTTTTGCG	GTCCGTAGAA	GATACCATTT	11640
TCTTTGTAAA	CTTTGATGGC	ATCAAGGTCG	CTCGGCATGT	TGGCACCTTC	AGATACACAG	11700
ATAACGCCTT	GAGCAACCAA	ACGTTTAGCT	GCTTCACCGT	TGATTTTCGTT	TTGAGTGGCA	11760
CATGGAAGAG	CAATGTCATA	GTTTCCAGCG	TAAGTCCATA	CAGTACCTTC	GTGGTAGGTT	11820
GCAGTTGCTT	TTTCAGCTGC	ATACTCAGTC	AAACGAGCAC	GACGTTTTTC	TTAACATCA	11880
ACCAAAAGAT	CGAAGTCGAT	ACCATTTTCA	TCGATGACAT	AACCATTTGA	GTCAGAAACA	11940
GAAATAACAG	TTGCACCGAG	TTCAGTTGCT	TTTTGAAGAG	CATATTGAGC	AACGTTACCA	12000
GAACCTGAAA	TAACGACTTT	CTTACCAGCA	AAGCTGTTAC	CGTTAGCTTT	GAGCATTTCT	12060
TCAGTATAGT	AAACCAAACC	GTAACCAGTT	GCTTCTGGAC	GAATCAAGCT	ACCACCAAAT	12120
CCAAGAGGTT	TACCAGTCAA	GACACCAGCA	TCAAATTGGT	TAAGACGTTT	GTATTGACCG	12180
TAAAGGTAAC	CAATTTACG	TCCACCAACA	CCGATATCAC	CAGCAGGTAC	GTCAAGTGAT	12240
GGTCCGATGT	GTTTTTGCAA	TPCAGTCATG	AAGCTTTGGC	AGAAGCGCAT	CACTTCAGCA	12300
TCTGTTTTAC	CTTTAGGATC	GAAGTCTGAT	CCACCTTTAC	CTCCACCGAT	AGGAAGTCCA	12360
GTCAGACAT	TTTTAAAGAT	TTGTTCAAAT	CCGAGGAATT	TCAAGATCCC	TTGGTTTACA	12420
GTTGGGTGGA	AACGAAGTCC	ACCTTTGTAT	GGTCCAACAG	CTGAGTTGAA	TTGAACACGG	12480
TAACCACGGT	TACTTGAAT	TTTTCCATCA	CGGTCAACCC	AAGGAACACG	GAAAGAAACC	12540
ACGCGCTCAG	GCTCAGTAAT	ACGTGCCAAG	ATATTTTCTT	CGATATACTC	AGGGTGTTTT	12600
TCAAATACAG	GTTCTAAAGT	GTTGAAAAAT	TCTTCAACAG	CTTGAGGAA	TTCAGCCTCG	12660
TGCCGG						12666

(2) INFORMATION FOR SEQ ID NO: 138:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 3083 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 138:

AGCAACTGTT	GTGAACCAAT	TCCGATAAAT	TCCAAGAATT	GGTTAATAGA	GCCATTTTGA	60
CCAAAAATCC	CGATAAAAGC	ATAGGCTTTA	AGGAGCAAAT	TGATCCAGGT	AGGAAGGATA	120
ATCAGCATGA	GCCAGAGTTG	ACGGTGTTTG	AGACGGGTCA	AAAAGAGGGC	CGTCGGATAA	180
CTGATAAGCA	GTGCCACAAA	GGTCACAATG	CCTGCATAAA	GCACTGAGTT	GAAACTCATT	240
TTAAGATAGG	TCAAGTTTGG	TGACGCAAAG	TAAGATTTGT	AATTTTCTAA	ACTGAACTGG	300
CCTTCGATGT	TGAAAAAGGA	TTGACCGAAA	ATCAAGACCA	AGGGTGCCAA	TACAAAGAGC	360
GCAATCCAAA	GCATGTAGGG	TACTACAAAAG	AGTTTAGAGC	TTGTTTTTCT	CATCTCTTTC	420
CTCCTCGATT	GCATTGATCA	AACCTGCTTC	TTGCTCTTCG	ATTTCTACGT	ACTCCTCAAT	480
ACGAGCATCG	AACTCTTCTT	CGGTTTCATT	GAGACGCATG	ATGTGGATGT	CTTCTGGTTC	540
AAAGTCCAGA	CCGATTTCCCT	CACCCACGAT	AGCCTTACGG	GTGAGTGGGA	TCATCCATTC	600
ATTTCCAAGT	TCGTCAATAG	CGATAAATTC	ATAATGAACT	CCACGGAAAA	GCTGGGTATC	660
GACCTTAACT	TGGAGCTTGC	CTTCTTCAGG	AAGGGTAATG	CGCAAGTCCT	CTGGACGAAT	720
AACGACCTCA	ACAGGTTTCAT	TTGGCTTCAT	CCCACCATCA	ACCGCTTCAA	AGCGTTTGCC	780
GTTAAATTCG	ACCAAGTAGT	CCTCAATCAT	GGTACCTGGC	AAGATGTTTG	ACTCCCGGAT	840
AAAGGTGGCA	ACAAAGTGGT	TGATTGGCTC	ATCGTAGATG	TCCACAGGGG	TCCAGACTG	900
GACAATCTCG	CCATCATTCA	TAACGAAAAAT	CCAGTCACTC	ATGGCAAGAG	CTTCTTCTCG	960
ATCGTGAGTG	ACAAAGACAA	AGGTAATGCC	CAATCGTTGT	TGTAATTCAC	GCAATTCGTA	1020
CTGCATGTCT	GTTCTCAATT	TCAAGTCCAG	CGCTGATAAA	GGCTCGTCCA	ACAAGACCAC	1080
ACGGGGTTGG	TTGATGATAG	CACGGGCGAT	GGCCACACGC	TGACGTTGTC	CTCCAGAAAG	1140
TTTGCGGATG	GAACGTTTTT	CATAACCTTC	CAACTGAACC	ATCTTGAGAA	CTTCCGCTAC	1200
ACGCTGCTCG	ATTTCTTTCT	TATCAATTTT	ACGCAAGCGA	AGTGAAAGG	CAACATTTTC	1260
AAACACATTC	ATATGTGGGA	ACAAGGCATA	GGATTGGAAG	ACGGTATGTA	CGTCGCGCTT	1320
GTTGGTTGGA	ATATCATTGA	TACGAACACC	GTCTAGCATG	ATATCTCCTG	TCGTGCGATC	1380
CAGTAAACCT	GCAATAATGT	TTAGGATAGT	TGATTTCCCC	GAACCAGATG	CACCTAGAAG	1440
GGTGTAGAAT	TTCCCTTCTT	CCAACATAAA	GTTGATGTCT	TTGAGAACCT	TGGTGTGCT	1500
GTCTTCAAAA	ACTTTAGAGA	CGTTTTTGAA	TTGCATAATT	GGCTTTTTC	ATTGGCATAA	1560
ATTCCTTCTT	TTTCATAGAT	TAACCGATCG	GGGCTCTGTC	AGGTCCCCAC	TACCTCTTGC	1620
AGGGAGTAAA	ACCACCTGCA	TACATCTTCG	CTACCGATAG	GCTTTCACCC	AAGATCCGGA	1680

930

CTTCTCTTTC	AAGCGTAATA	CCTGAGTGTT	CCTTGACTTT	TTCGATAACC	GATTGGATCA	1740
AGTCCTCGTA	GTCTTTGGCC	GTTCCATCTG	CGACATTGAT	CATAAATCCT	GCATGCTTTT	1800
CTGACACTTC	TACGCCACCG	ATACGATAGC	CTTTCAAGCC	AGCTTCTGAA	ATTAAGTAC	1860
CTGCAAAATG	CCCGACTGGA	CGCTTAAAGA	CCGAGCCACA	AGATGGGTAT	TCCAAAGGTT	1920
GCTTGAGTTC	ACGTAGGTGC	GTCAAGCGGT	CCATTTCCCTG	CTTGATAACC	TGATGGGTTC	1980
CTGGAGCTAG	GGCAAATTTA	ACTGACAAGA	CAACTGCACC	AGACTCCTGA	ATAGCTGAAT	2040
GACGGTAACC	AAAAGCCAAG	TCTTTAGCAG	ACAGGGTTTC	GATTTCTCCA	TCCTTGGTCA	2100
AGACCTTACA	AGACTGCAAG	ATGTGAGCAA	TCTCGCCACC	ATAGGCACCC	GCATTCATAA	2160
AGACAGCACC	GCCAACGCTT	CCTGGAATAC	CACAAGCAAA	CTCAAAGCCA	GTTAAACTAT	2220
GACGGAGGGC	AATGCGAGTT	GTTTCAATCA	AGTTAGCCCC	AGCTTCTGCT	TCAATGGTAT	2280
AGCCATCAAC	AGAAACGTTA	TTGAGCTTGT	CACACAAGAT	GACAAATCCA	CGAATCCCAC	2340
CATCACGAAC	GATGATATTG	CTTGCATTGC	CAAGAACCAT	CCAAGGGATA	TTTTCTTGGT	2400
TGGCAAATTT	CACAACGCGA	GCCAACCTAA	AACGATTTTCG	TGGAAAGACC	AAATAATCAG	2460
CCTCTCCACC	TACTTTTGTA	TAAGTATAGC	TATGCAAGGG	TTCCTTAAAA	CGGATATCAA	2520
TTCCCTTCTAA	GATTTCAAGC	ATTTTCTCTC	TTACAGACAT	GTCACTCTTC	CTTTTACAAA	2580
ATTCATTCCA	TTATACCATT	TTTAGAGACA	TTTGACGACC	ATAAAAATAC	CTTGTTTGGG	2640
TTTTGCATAA	GAAAAAGAGG	TTCCCCCCTT	TTTATGATTT	TTTACAAAAG	ATTTCCCTGG	2700
TTCCATAGGC	GACCAGAACG	AGCTCCAGTG	CTAGAATCAC	TTCAACCAAG	ACTGGATTTG	2760
TCAACCAGCC	TACTTGGAAG	AGAGATGGTG	CCAGATCAAA	GAAGGCATGC	AAGCCATAGG	2820
CTGCTAGGAG	ATAAATCCAT	TTCTTCTGGC	GAACAGCTTG	GTA AACCCAA	ACTGTCAAAA	2880
GTAATTTGGAA	ACCAAGCGCC	AAGATTCGCT	CAAAACCAAG	CAAATAAATC	TGCCAGACCG	2940
AAAGTGAAGT	AATGGTTTTT	AACATATTTT	CAGACAGTAA	TTGCATAACC	TGTGGATTC	3000
GAGTTTGAAC	TGCCGAAAGA	ACAATGTAAA	GATGAGTAA	ACTAGTAAGG	CCTAGAAAAA	3060
TCAACTCCAA	GCCACCATGC	CCC				3083

(2) INFORMATION FOR SEQ ID NO: 139:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15363 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 139:

CCGGAGGATA	TTGACCACCA	CCAAAAGCAG	GGGGAAAATC	GAAATCAACC	AATAGTAGGC	60
TACTGCGACA	CTGGTCAACT	CACTATCTGA	TGCTTGATAA	TAATGCAAAA	AAGCTTTTAA	120
TAAAGGTTTG	TCTATCAGCT	CTTTCACCA	CTTTTTCATG	TCATACTCCT	TCACTTATAA	180
TCTTATACTC	AATGAAAATC	AAAGAGCAAA	CTAGAAAGCT	AGCCGCAAGC	TGCTCAAAAC	240
ACTGTTTTGA	GGTTGTAGAT	AAGACTGACG	AAGTCGATCA	CATACATACG	GTAAGGCGAC	300
GCTGACGTGG	TTTGAAGAGA	TTTTCGAAGA	GTATTAACTA	ATTTCTTCTT	ACCAATTCCA	360
CCATATCATA	CGGTAGGGTA	TTGGCAGCTT	CCTTCAAGGA	ATAGTTCTCT	AAGTTATTTA	420
CATTTTGTGC	TAATTTCTTG	GCATACTTAG	TCGTAATCAA	TCGTTTTTCT	TCGTATTCGA	480
AAATCAACTT	GCGCTCCAGA	TAATAGCCTC	TCAGCATTTC	ATCGATATTG	TGGGTTTGA	540
CACGATTGAT	AACCCGTTTC	ACAAAGGCAC	CACTGCTGAT	AATAGCTGTT	TCTCGAAGAC	600
GAGACTCCTG	CATAAACTA	ATCAAAGAGC	GTCTGTAGAC	TCCCTTCAGG	TTTTCCAAC	660
TTTCAATAAT	CATCTCTGTA	TTGGCAAGAT	AGAGCTCTGC	AATTTGGTCA	TAATCAAGAG	720
CACGGAGACG	GCTTTGCTCC	TTGTTCTTCC	AGCTACGGAA	GGTCTTCCG	AGAGTAAAAA	780
CTTCATGAAG	GAGAAAACGT	AAAATCCTCA	AGGAAACAAG	AAAATAATAG	GTCAGTCTTG	840
AGGCAAGTTT	ACGATTGATT	CCTTGTCTA	TATTTTTCAG	ATAACGTTGG	TAAACTCGGT	900
AAGCACGATT	GCTAATGTTT	CCCTCTTCAT	AGGCCTGTTT	CAAACCATCA	CTTTCAATAC	960
TAAGAATCAA	GAGTTTCAAA	GCAGCCCAGT	CTTCTTGATC	ATCCTGGTTT	TCTTGGCTTA	1020
AAATGAGATT	TTCAATACGT	CCATGATAAT	TGTCAATAGC	CGCATAGAGG	GGAAGTTTAT	1080
TTCTGGTGTC	TTCCAACCTT	TTTTCCAACCT	CTAGCGTTAC	TTCATTCAAA	ATGGCGATAT	1140
GCATAAGATA	ATCCTTGCTT	TCTTCTCTTT	CATCAGAAAG	ATGAGGCAAG	ACCAAGAGAC	1200
CTGTTAAAAA	GCTAACCAAGC	GTCACACCTG	CAACAAGGAA	AAGCAAAAAG	GGATACTCCT	1260
GTTCTAGATT	ACTTGGTATC	AAGAGAATCG	TAGCAATCGA	CACCGTTCCC	TTAACACCTG	1320
AAAAGGTCAA	GAGAAACATG	TCCTTCATAT	ACTTATTTAG	CTTTTCTTTG	AGGCGTCGGG	1380
TTCTATAGGC	ATAATAGCCA	TAGATCATAA	TAAAACGAAT	GACAAAAAGG	ACAAAGGTAA	1440
GGGCGATAAG	AGATAGCAAT	AAAAGTAGAG	GATTATAGAT	TGGATTGGTC	AAGATAGGTT	1500
CTGCTATCAT	TTCCAACCTC	ATCCCTAAAA	TCACAAAGAC	AGAACCCTTG	AGCATAAAGG	1560
TCACTGTATG	CCAGACCGTC	TCGGTCACCG	TATCCACTTG	GGCTTCGAGG	AGCGTGATTT	1620
TCTTGAAGCG	ACTTGCTTTT	AAAATTCAG	CAACTACGAC	GGCAATAATA	CCTGAAACAT	1680
GAACTTCTTC	TGCCAGAAAG	AAGGTCACTA	GAGGCAAACT	CAATTCTAAT	AAAAGTTCAC	1740

TGGCAATATC	CGTTGCGCGC	ACACTTAGCA	AGAAGGTATG	GAGGAAGCGG	TTGGTCATGG	1800
CTGTTAAAAA	TCCAATTA	AAACCGCCTA	GGATTGAAAA	GATGAGCGAA	CTGCTAGCTT	1860
GCCCCAGAGA	AAAAGCTCCA	GTTGTCCAAG	CTGTCAAAGC	TACCTGAAAA	GCCACCAAAC	1920
CAGAAGCATC	ATTCAAGAGT	CCTTCGCCCT	TAAGAATATT	GGACACGCGC	TTAGGAAAGC	1980
TAAAACGCTC	CGAAAGAGAG	GCAAAGGCCA	CCAAGTCCGT	AGGACCAAGG	GCTGCCCAA	2040
CAGCCAAGCA	AGCTGCCAAG	GGAAGGCTGA	ACCAAAGAAG	ATGGGCCAAG	CCACCAAAC	2100
TCAGGGTCGA	GATAAAAAATC	ACTGGAAATA	TGAGATAAAC	AATGATTCGC	CAGTGTTTTA	2160
AAATAGCCGT	AACATCTGCT	TCTTCAGCCT	CTCGAAAAAG	CAAGGGTCCG	ATAACCAAGT	2220
CCAAAAACAA	CTCCGTATTA	AGGTGAAAGT	CAGTATTGGG	TAAAAAGAGA	CCAATCACAA	2280
TTCCCAAAG	AATTTGCACC	AAAGGGAGAG	GCAAAAAGGG	CAGGAGCTTA	TTGGTTGTAC	2340
TTGAGACAAT	CAAAACCAGT	AAAAATAGGA	TGAGGTAAAT	CAGTAATTCC	ACGCACGTCC	2400
TCCTTAATCT	TTTTTACAAC	AGGATTCAAA	TATCTCCTTC	TGCTCTTTGA	TTTTTTGGTC	2460
AATCTTGGA	CAGTCTTTGT	GCTCAATTTT	TCTCTGGCAC	CGTTCCATTT	CAAGAGCAAC	2520
TAATTTTTTC	TTGATTTTAA	GCATTTTTTT	GCTCATATGC	GCTTGGTCTA	GCACGCCCAT	2580
CGCTCGTTGC	TGGTGGGTTG	ATTCAACAAA	ATTCTGGCGC	ATGGCATCCA	GCTTTTCGTG	2640
TAAGTATTGT	TTATCCATGT	CTGTATCTCT	CTAATTTTTC	AATCATCACT	AAAAACGGCG	2700
GGTTGTGAC	TTGGTTTAAA	GTTCCGTAAA	TGGCAGCTGT	GTACTCTTGT	TGGTTCAACT	2760
GGATCACAAA	ATCCAAGACA	GCATCTCTCT	CGAGATCGCC	TCCTTCATGA	CCATAGTAAA	2820
TCATAATAGC	AATTCGTCCA	CCTTTGACAA	GTAAGCCACA	TAGCTTTTCT	AATGCCTCAA	2880
TCGTTGTCTG	CGGTCGGGTG	ATGACAGACT	TATCAGCTGC	CGGCAAATAG	CCCAGATTAA	2940
AAATCCCTGC	CTTAGCTTTT	ATCACAAACT	GGTCCAGTGT	CTCATGGCCT	TGCAAGATTA	3000
ACTGGGCATT	TGTCAAGTCA	GCCTGATGCA	AACGCTCTTG	GGTCTTTTCC	AAGGCTTGCT	3060
TCTGAATATC	AAAGGCATAG	ACTTGCTTGG	CTAGCTTGGC	TAAAAAAAGC	GTGTCATGAC	3120
CATTTCCCAT	AGTCGCATCC	ACTACGACAT	CCTCTTTTGT	CACGACCTCA	GCCAAAAAAT	3180
CATGTCCCAT	CTCAAGTGGT	CTTTTCATTT	TCAAACCTCT	GTTTTACAGC	CTTGCATCCT	3240
TGAACACTTC	CACGACGTCG	CATCTCCATC	TCAATGCTGT	TGAGGACTTC	CCATTTATTG	3300
AGGCTCCACA	TAGGACCAAG	CAGCATATCC	CTAGGCGCAT	CTCCTGTAAT	TCGATGGATG	3360
ACGATATGTT	TGGGAATAAT	TTCCAGTTGG	TCACAGATGA	CCCTGACATA	TTTCCTCTGA	3420
CTCATCAATT	GTAACGCCCC	CTCATGGTAA	TCTCGTTGCA	TACGAGTATT	TGTCATAAGA	3480
TGGAGCAAA	GCAGTTTAA	CCCTTGAATA	TCGTTATCCG	TGACACAACG	GCGGACATTT	3540

TCAACCATCA	TCTCATGGGT	TTCACCAGGC	AAACCATTGA	TCAAATGGGA	AACAATCTCA	3600
ATTTTGGAT	ACTTCTCAA	ACGCTTGACC	GTTTCCACCT	ACAATTCATA	AGAATGCGCA	3660
CGGTTAATCA	GGTCAGAGGT	TGCTTCATAA	GTAGTTTGCA	AGCCCAATTC	AACCGTCACA	3720
TGCATGCACT	CCGATAACTC	AGCCAAATAT	TCGATGGTTT	CGTCTGGTAA	ACAGTCTGGG	3780
CGCGTTCCAA	TATTGATTCC	TACCACACCT	GGCTCATTGA	TAGCCTGTTC	ATAACGCTCT	3840
CGAATAAATT	CCACCTTTTC	ATGGGTGTTG	GTAAATTTT	GAAAATAAAC	CAGATACTTC	3900
CGAACATCCG	GCCACTTGCG	GTGCATAAAG	TCAATTTCCCT	TATAAAATTG	CTCACGGATA	3960
GGCGCATCCG	GTGCCACAAT	GGCATCTCCA	GAACCAGAAA	CCGTACAAAA	AGTACAGCCC	4020
CCATGAGCCA	CAGTCCCATC	ACGATTGGGA	CAATCAAATC	CCGCATCAAT	AGGGACTTTA	4080
AAAGTCTTTT	CTCCAAAGAG	TTTTCGATAA	TAATCATTCA	AGGTATTATA	AGATTTCATG	4140
ACTTTCATTA	TAACAAAAAT	CACCCACAAT	CTCAAAAGCC	TGACTTTCCT	ATAAATTCCT	4200
CTGTTTCTCG	TTTCCATTAG	CCTTTTTHA	TGATACAATA	TGGGTATGAT	TTAATGAAA	4260
TTAGCATCTA	TTTTATTATT	GATACTGACC	TTAGTCGTCT	GCATTATCCT	AACCAAATT	4320
TTTAGATTAA	AAAACTAGG	ACGAACTTT	GCGGATTTGG	CTTTTCCAGT	CTTGGTATTT	4380
GAGTATTACT	TGATTACAGC	TAAAACCTTT	ACCCATAATT	TCCTCCCTAG	ACTGGGGCTA	4440
GCCCTCTCGA	TCCTAGCCAT	TATCTCTGTC	TTTTTCTTCC	TTTTGAAAAA	ACGCAGCTTT	4500
TACTACCCTA	AATTTATCAA	ATTCTTCTGG	CGTGCAGGAT	TCTTATTAAC	CCTTATCATG	4560
TATATAGAAA	TGATTGTTGA	ATTGTCTTA	ATGAAATAGT	CGAATCCCTA	AGCATTTTCT	4620
AGGGATTTTT	GCTTCTCTA	CAAAATAGTA	TAGACAATAA	CACTATACAA	TTTTATACAA	4680
AGAAAAGAGT	CTGGGACAAT	AGTCTCTTAT	ATCCAAAAAG	GCAACGGATT	TGCCGTTGCT	4740
TTTTTGGATG	GTTACGATAG	TCTTGGTAAA	ATAGAATTGC	CCAATAAACC	ATTTAGAAAAG	4800
GCTATCCCAT	GCATATTCAC	TATAACACAA	ATCAAACAAC	TTTACCACTA	GAAATCAGTT	4860
CCTTCTTACC	ACAAGATCAT	CTCGTTTTTA	CTATTGAAAA	AGTGGTGAAT	ACCTTGAGG	4920
AACGTCACCT	CTACACCTCC	TATCATGCCT	TTGATCGCCC	GTCTTATCAC	CCTAAAATGC	4980
TTGTATCTAC	TCTTCTATTT	GCCTATTCAC	AAGGGATTTT	CTCTGGTCGA	AAAATTGAAA	5040
AATGGAAGAG	TTAGTGACCT	TAGATTGTTT	GTTTATTGAC	AGAACTAAGA	TTGAAGCCAA	5100
TGCCAACAAG	TATAGTTTTG	TGTGGAAGAA	AACGACAGAG	AAATCTCCG	CCAACTTCA	5160
AGAACAGATA	CAGGTCTATT	TTCAAGAAGA	AATCACTCCC	CTTCTGATTA	AATATGCCAT	5220
GTTTGATAAG	AAACAAAAGA	GAGGGTATAA	AGAGTCAGCT	AAAACTTAG	CGAATTGGCA	5280

CTATAATGAC	AAGGAGGATA	GCTACACACA	TCCTGATGGC	TGGTATTATC	GTTTTACCA	5340
TACCAAATAT	CAGAAAACAC	AGACAGACTT	TCAACAAGAA	ATCAAGGTTT	ACTACGCCGA	5400
CGAACCTGAA	TCAGCCCCTC	AAAAGGGACT	GTATATGAAC	GAACGCTATC	AAAAC TTGAA	5460
AGCTAAAGAA	TGTCAGGCGC	T TTTATCTCC	CCAAGGTAGA	CAGATTTTCG	CTCAACGCAA	5520
GATTGATGTG	GAACCTGTCT	TTGGGCAGAT	AAAGGCTTCT	TTGGGT TACA	AGAGATGTAA	5580
TCTGAGAGGG	AAGCGTCAAG	TGAGAATTGA	CATGGGATTG	G TACTTATGG	CCAATAACCT	5640
CCTAAAATAT	AGTAAAATGA	AATAAGAACA	GGACAAATCG	ATAAGGACAA	TCAAATCGAT	5700
TTCTAACAAAT	GTTTTAGAAG	TAAAAGTGTA	CTATTCTAGT	TTCAATCTAC	TATACAATAA	5760
GAGAA TGACT	CAAAAT TAAA	AAGCTAGAGT	TCCACAATTG	GAAATATCTA	GCTTTTTTGT	5820
GGTTGAGAAC	TATTTTGTCT	CAGGCTCTTT	ATCTTCTATT	TAGGACAAGA	GTTTTTCTTT	5880
GGTCTTTAAT	GATAAAGAAG	GTATCAAAAT	TTCTAGTCTT	CTTTTTTACC	TTTAGTAACT	5940
ACTAATCCTG	CACTCAAACC	TAGAAGAGTT	AAACCTGCTG	CTACTGCTGC	TTGGCTTGCC	6000
GCACTACCTG	TACTTGGTAA	CTGGGCTTTA	TTAGTTTGAC	TAGCTTCACT	TGAATCAATT	6060
GGTTTTGTAT	CTGCTTTTTC	TGACACTTGT	GGTTTTTTAG	CTTCTTGAGC	TACTGGTTTG	6120
GTTCCAACCA	AGACGATGCG	GTCTGTCCGA	ACTTCTACCA	CTTCACGGAG	TTTTTCTTCC	6180
TTACTTCCAT	CAGGATTAAT	CGCTGTAAAG	ATACGTCTTT	TTCCAAC TTT	TCCTTCTTGT	6240
TCTACACGAG	TTTCACCTAG	ATACAGTGTT	GAATCTTTTT	TCTCAACTGT	CTTGTATGCC	6300
AAATCTTTTT	CAACAAATTC	GATTTTTTGA	AGATCTTCTT	GTACAGCAGC	AACTGTCTTC	6360
TCAGAAACTG	GTTTTTCCTT	AGTCAAGTGG	ATACGGTATT	CCTTGACTTG	TTTTCCACTT	6420
TCTGAAACGA	GGCGAACAAAG	TACTGGAAAG	CTATCTTCTC	CACTATCTAC	CACAGTTGAA	6480
GCTACTTGAT	TGTTTTCTTC	AACTGAGACT	TTTGCCGTT	GACCTT TATA	GGTAATT TGA	6540
TAGTCTTGAC	GATTTTCAGC	GAAATCAGCA	AGTTCTTTTC	CATCTACAAG	AATCTTTGAT	6600
TGAGTGCTTT	CTTGAGGCAA	TTCACTTGGT	GCAAGGAAGG	TCATCTCAAT	CATCGCAACA	6660
CCGCTCTTAT	CTGCTTTACG	CTCCATACGC	CATCTCATAG	CTTTGGCTTT	GATAGCTTTA	6720
AATGTTACGT	TGATTTTATC	ACCAGCTGCA	ATGTCTTTAT	CCGCACGATA	AGGAACAGCT	6780
TCCCAATTTT	CTGGATTGTT	GAATGGATGG	TCTGCGTCGT	AGGCTTGGTA	GTTTGAATAG	6840
TAGGTTGGCA	CTTCAAATC	TGGACCGACA	TAGCGTTCTA	AAACGACTTT	AGATGGTGCA	6900
TCCGTACCAC	TATCTGCAAA	GAAC TGAAC T	TTTCC TTGTG	TAACAGTCCG	TTCTACAATC	6960
TTACCATTTT	CACGGAAAAT	CACACCCGCT	GATACTTCTG	GATTAGAAGA	TGGTGT TGGT	7020
GACCAGTTTG	TCCAACGACG	ATTTTCTGAA	TGATCTCCGT	CATTGAGATA	GTCAACGCGG	7080

TCATGAGAGT	TTTTGTCAAT	ATCATTGGTT	GCTGAAGCAA	AGGCCTGGTT	ACTGTTTTCA	7140
TCATAGTTAG	GGTTATCTGA	AAGAGTCTCA	CCAAGTTTGT	CTGTCACTCG	TACAGTGATC	7200
TCAGCAACAA	GGTACTACC	AAGGACACGG	CCTCGAACAG	TAAATTGACC	TGCTTTTGTC	7260
AGATTTTCCG	CTGGAAC TTC	TTCCCATTC	ACTGTCAGGT	CTTTTGTTC	GTAGCCGTCT	7320
TTACCTGTGA	AGTAAACTGG	AACCTTAGTC	GGCAATTCAA	GTGCTTGACC	TACTTGTAGC	7380
AAGCGAGCTT	GTTTAACCGC	AGCAACTGGT	TTATGAGAAA	GTAAGCTCTT	ATCCTTAGTG	7440
AAGTGCAGAC	GGTATTCTCC	TAAGATGTCG	CCATTTTCAG	CTTTCGCGAT	GACACGAACT	7500
GGCTCACCTT	CACGAACGCT	TGGAACGACG	GTAGCGAGAC	CATTGTTGCT	AACACTTGCT	7560
GTGACTGCCG	GAAC TTTTCC	ATCTACAGAC	TCAAGGTAGT	AGTCTGTCAA	ATCAGGGTTG	7620
AAGTTTGCTA	AGTCTTTGCC	GTCAACTTGG	ATTCTTGT TT	GTCCTTGCTT	GGCTGCCGCA	7680
ACTTGTTCG	CAAAGATT TG	TACCTCTGTG	ATAGACG TTC	CACGCTTGTT	ATCTGCTTTA	7740
ACCATGCGAA	TACGAACAGC	ATAGGTTTCA	ACTTATCAA	AGCTAAAGTG	GTTCA TTTCT	7800
CCAGCCTTGA	GTTGAGCAGG	GGCTTTTAGA	TTAGTAAC TG	GTTTCCAGTT	GGCAGAATCA	7860
TTAAAGACAT	GGTCCTCAT T	ACCAACAAAA	CTAGGGTTT T	TAGGAGCTGT	TGGGACAGTC	7920
TTACCAACAT	AATACTCAAT	CACATAAGAC	TTCGGTACAC	CAACTCCATG	GTCTTCATGG	7980
AATCCGACAC	TTAGATTATC	AACGGAGCGT	TTGCTCAAGA	TACCTGAATC	TCCAAACAGA	8040
ACACCGACTG	AAGCTTCTGG	ATTAGTACGA	TTCCAGTTTG	TCCAACGATT	GGCTGGTTGG	8100
TTATGTAGG	AAATGAGCTT	GTCATTAACA	TTTGAAC TG	GGTCGCTTGG	ATTTGAGTCT	8160
GAAGCAAAGG	CAAGTGGCAA	TTCTGAACCG	GTCCATTGGT	CAGAAATGTT	TGCACCTTGC	8220
TCAGTTTGAG	CAGATACGCG	AACATGAAGT	TTAGTTGTTA	ATTGCGTACC	TTCTAAGCGA	8280
CCATTAAC TG	TAAAGACACC	TTCTTAGCG	TATTGCTCTG	GACGAATCGC	ATCCCATGCA	8340
ACCTTAGCTG	ATGAAACG TG	ACCATTTGAA	TCATATGTCC	GAACACTTTC	TGGTAATTGT	8400
GGTGCTTCTG	CGATTGGAGT	TGTCACACTG	ACTTCTTCAA	CTGAAACGAT	ACCTTCTACA	8460
GAGACTTTTG	CACGCGCTTC	AAGGTCAAT T	CCTTCAACTT	TACCTAGTAC	TTCAAATGTT	8520
TGATAGGAGT	CTAGTTTTTC	TTTCGGAATA	GCTTGCCAAG	TGACTTTATG	AGTTT TAGGG	8580
AAACCTTTGT	CATACTCAAC	TGTTACTGTT	GCTGGAAGAC	TTGGTTCC TG	ATGCAAATCT	8640
GTCACTACAT	TTACAGGACG	GATGGATTGC	GCAATCTTCT	TCTCAGTATT	GGCTTG GATA	8700
GTGAGTTCAA	CTTGGTCTTT	AGCTCCCTCA	TATTCAGCGT	TCAGAGTGAC	TGCTCCTGGC	8760
TTATGCAACT	CAAGCATTCC	TTTACGAAT T	GCGACTTCCC	CTTCACCACT	TGTAGAGAAG	8820

GTTACTTTAT CAGCTGGTAA TACAGCTTGC GTTCCATCTT GATAGTGAGC TCGAACCGAC 8880
 AATTTGACAG TTTGGTCTTC TTTGAGACTG TCAGCTTTTT CCACTTGCAA GCTCAAGTGA 8940
 GCAATTTTTG GCGCTTCTTC AAGGAATTGA ATTGCATAGG TTTGAAGAGG GCCACCATCT 9000
 TTAGGCTGAA TAAAGATGCT CGCACGCATG CCGTTTGCTG CGCTTGCTTG AAGAACTGTA 9060
 ACAGCTGCAT TTTTAGCACT TGCTGTGACT TCTGGCAACT TAGCTCCATA AGCAAGAGTG 9120
 CGGTATTGCA TTGGTTTTTG ACTAGTAAGA CCTGTACTG CCTCACCACC AACCGTTACA 9180
 GTTGGTACTG CAGGTGCCGC AGGATTGCCT TCTTCTACCA CAAGGGTTC ATGAATTGGT 9240
 TGACCTCTA AATAACCGGT CGCTTGAATA CGAGAACCTG GAATTGCTAA CTTAGCTTTA 9300
 TCTTCTTCGG CAATCTCCCA CTGTGCCACT TCATACTCTT CAACACTTCC ATCAATCAAA 9360
 ACATAGGAAA CAGATTTGTC TACAGAATTC AAGTCAGTAT TTGGAGCAAT ACGTTTCACA 9420
 ACTGGTAGCT CTGATTTAAG AGCAATCACT TCTACACGAG CTTCTACTTC TCGTCCGTCA 9480
 GCCATACCTT TCACCGTTAC AATACCAGGC TTGCTCACAT CTAAGAAGA CCAGGTTACA 9540
 GGACGTTCTG CACGGCTACC ATCACTGTAT ACAAACGGAA CAGTGGTAGG CATTTCAGGT 9600
 GCCTCTCAA TAATGGTCTG TACTTTTGGC ACTTCTGTCC CAAAACAGT CTTCTCTTGT 9660
 CCTTCTTCT TACCAGTAAA GACAGTGACT TGTTTCGATT TCAAGAGATC AGAGTGGGCA 9720
 GTCAGGGTGA ATTTCCCTGC TTGTTTCAGTT GATTTGACAA TGGCAACACC TTTACCATTA 9780
 AATGCTTTAC GAATCCAAGA ACCATCTGCT TGCGCCTTAT AGCGTTCACG GCTGGCTTGT 9840
 TCTCCGTAT CTACACCGAC CAGTTGACCT TGCCCATGCA ATTGGAAGCG AACCAGATTA 9900
 TTAGCAGTTG GAACCACATT CCCCTGGCTG TCAACAATTT CATAGTAGAT GTAAGTCAAG 9960
 TCTTTTCCAT CTGCTGCAAT CGCATGGTCT TCCTTAATAA GACGAACTGC CGCTGGCTTA 10020
 CCAGCAGTCG TAATCTTATC TCGAGCAATT TCCTTGCCAG ATTCATCACG AGCAATTGCT 10080
 TCCAAGTAC CTGGTTGATA GGCAACTTTC CATTCAAGAT AAAGTTCATT AGCATTTGCA 10140
 CCTTCTTGGT AAGTCCGCC ATCGCTGGTT TGTTTTTTAT TGAAAGTCTT AAGACCAAGA 10200
 GATTTTCCAT TCAAGAACAA TTCTACACTA GAAGCATTCG AATAAGCACG AACTGGAATC 10260
 TTACCTTCTG AGTCAGCTAC TTTGGATGCT AATTCTTTGT TTTCCAGTT CCAGTGAGGA 10320
 AGAAGGTGTA CCATCGGTTT CTTCTTAACA GAAACCCATT GGCTTTGGTA GAGATAGAAG 10380
 TCATGTTTTG GAATGCCGGC TGTATCTACG ATACCAAAGT AAGAGCTCTT AACAGGAGTT 10440
 TGATTTTGGT TGTGCCATGG TGTAGGTTCA CCAATATAGT CCGTACCTGT CCAGATAAAC 10500
 TGTCCAGCAT AGCCAGCGTT GTCACGGTCA AAAGTCCATG AAGCGGTTCG TGTTTTCCCC 10560
 CAACCCACAC GATCATTTCC ATAATCTGAC TGTTTATAAT TACGCTCAGG TCCATTGCTA 10620

TGTTTCAATT	CACG TTCAGG	GCGATAGTAA	CTTCCACGTG	TACGGGTAGC	TGAAGATGTT	10680
TCTGATCCAT	AAATCAACCA	TTTTGGATGC	TTAGCTCTAA	GGGCTTTGTA	ATTATCTTCA	10740
GAATAGTTAA	ATCCAACAGC	ATCGAGTTCA	TCAGCAATTT	TCTCATGCCC	TCCGCTACCA	10800
TTACCGAAAC	GGAATTTATC	TGCTCCCATG	GTAACATAGC	GAGTCTTATC	AACATCCTTG	10860
ATAACCTTAA	CCAAACGTTT	AACAGTTGCT	AAAGAGTGGG	CATCACCATT	AGCTTCACCT	10920
ATTCATTAC	CAATTGACCA	CATGAAGATA	GCAGGGTGT	TTTTGCCTCT	TTCGACCATG	10980
GTACGTAGGT	CAAAATCAGA	CCATTTTTC	CCTTTTCGAG	CTTCTGGGTG	AGTGGCATCT	11040
TTTTCAAAGA	AACGTCCATA	GTCATAAGGT	TTCTTGCCAC	CATACCACGT	ATCAAAGGCC	11100
TCTTCCTGAA	CGAGTAAACC	TAGTTCGCT	GCGATTTGCA	AGGTTTGCTC	ACTAGCAGGG	11160
TTGTGGGTTG	TACGGATGGA	GTTAACTCCC	ATCTCCTTCA	TTTGTTTGAG	ACGGCGATAT	11220
TCTGCTTTAT	AGTTTTCTTC	TGCTCCAAGC	GCCCCATGGT	CGTGGTGCAA	GGATACTCCA	11280
TGGAATTTAA	TACG TTCACC	ATTCAAAGAG	AAACCTTCAT	TTGGAGTCCA	GTGATAGTAA	11340
CGGTAACCAA	ACAAATCCTT	CTTAGCATCA	ACCAATTGAC	CGTCACGGTA	AACACGCGTA	11400
ATCAATTCGT	ACAAGGCAGG	TTTGTCAATTT	AAAACAGTCC	AGAGTTTGG	TCTTTCAACT	11460
TCTAAATCG	CATCTAGGCT	TGTTGATTCA	TGTGCTTTTA	AGGTACGACT	CGCTGTACGA	11520
ACTAAGCCTG	TTACAGCATG	ACCACCTCGT	TCAACGATTT	GATATTCGGC	TACAAGTTCA	11580
TGGTCTTTGT	CGTCCGTATT	GACGATTTTG	CTGGTCACAT	GAGTTTCAAC	CTTGCCATGT	11640
TGTGTCTCTT	CAAGTTTGG	TGTTAAAATA	GTTGTCCCAT	TTTTCTCAAC	ATGCACCTTA	11700
TCTGTCACCT	GTAAGTCAC	ATCACGATAG	ATACCACTTC	CTGAATACCA	ACGGCTACTT	11760
GGCTGTTTGT	TGACTGCATG	GACAGCAATC	ACATCTCAC	GACCATCTTT	TTGAAGGTAT	11820
TTGGTGATAT	CATATGAGAA	CTGGTTATAA	CCATTTGGAT	AATGCCCCAC	TAAGTACCA	11880
TTGACATAAA	CTTGAGAATC	CATGTAGACG	CCATCAAAG	TAAGGCGAAC	ATTTTCTTTG	11940
AGGTCTTTTT	CATCTAGTTT	GAAAGTCTTG	CGATACCAAG	CTTCCCCACC	GTTGAGCTGT	12000
CCACCTTCAT	TTTGTGCAGG	AGATTCATGA	TCGAAATCGT	TAAAGATACT	CCAGTCATAC	12060
GGTAAATCTA	ATTTTTC	CGTAGATACG	TCTGCATCAG	GTTTAATGGC	TTCCCTAGAA	12120
TTTGCATTGA	GTTTAAAGTA	CCAATTTTGA	TTAAAATCCA	CTTTCCTGTC	TTCAATCATT	12180
TGATTCACCT	CTTCATTTGT	TACAGCTTTA	GCATCTTCCT	TGAGCGGTTT	TTCTTGATTT	12240
GAAGCTTGTG	ATTCATCCT	TGGAGCTTTT	TCTTCCGGTT	TAGCAGACAC	TTTTTCTCT	12300
TTTGGAGTTA	CGGCTTCATC	TTCTTCTTC	TCAGATGCAA	TAGCCTCAGT	TGAACTAGGT	12360

TCACCTTGTT	CTGTCCTTC	AACTATATTT	TTAGTTTCCA	AAGCTTTATC	AGCCTTTTCT	12420
TCTACTATCA	TTTTTTCCTC	TTTAGGTTTC	TCAGCAGTAT	GAGTAATAAG	TGTTTCATCC	12480
GCATAAACTA	CAGATTCTCC	AGCTATATTT	CCTCCTAATA	AAACTGCACA	AGTCCCAATC	12540
ATTACTGAGC	AAGCTCCAC	AGCAAACCTA	CGAATGCTAT	AAACTCTTTT	CCGATTCCAA	12600
TGGCCTTTCC	CCATAAAACC	CTCCTTATAT	TATATTTAGT	GCAGTTAGCT	ACTACCAAAG	12660
CCCAAGTGGT	ATACATGGTA	TGACAACCTA	GTTTCAACAA	TTTACTCTCT	GCGAAAATCC	12720
AATTCAAACT	TCGTCAGTGT	CGCCTTGCCG	TAGATATGAT	TACTGACTTC	GTCAGTTTCA	12780
TCTACAACCT	CAAACCATG	TTTTGAGCTG	ACTTCGTCAG	TTTCATCTAC	AACCTCAAAA	12840
CCATGTTTTG	AGCTGACTTC	GTCAGTTTCA	TCTACAACCT	CAAACCATG	TTTTGAGCTG	12900
ACTTCGTCAG	TCTTATCTAC	AACCTCAAAA	CTGTGTTTTG	AGCAACCTGC	GGCTAGCTTC	12960
CTAGTTTGCT	CTTTGATTTT	CATTGAGTTT	ATATTTTATA	GGAGCGCATT	ATTTTGCTTT	13020
TGCTGCGTAC	TCTTCGTTAC	GTTTGATCAT	TTGTTTTCTG	TACCAAGCAA	AGATACCGAT	13080
ATAGAATACA	AGGAAGACTA	CTGCACCAAG	GATTGCTTTG	ATATCACCAG	TTGTAGTGTT	13140
ACCAATTGTC	CAACCAAGAA	GTTTTTCGAT	TGGTCTTCA	AGAGTAGAGT	GAGTAATCAA	13200
TTGAGTTTGG	CTCACACCTT	CTGGGAAGGC	ACCTACACCT	TTAGCAAGTT	CTGTTGCAAA	13260
TGGTGCAATA	AGTGACCTG	AAAGAAGGAA	GAGTGGCAAC	AAGAGTGTTT	CGAAGATAAT	13320
CATACGGAGC	AATTTACCAC	GAGTTACAAC	CAAGAGAGCT	GGAGTAACAC	CCATAGCGAT	13380
GATACCTGCA	AGTGCAAGA	TACCATTTCC	AACTTTTGAA	AGAAGCACTG	CTTCAATCAA	13440
CATGATTGGT	GCAAGTACGT	TGGCACAAGC	CCAGATTTCA	GCACGACCAG	CGATGAATGG	13500
CCAGTCAAGA	CCGATATTGA	ATTTACGTCC	TTGAAGACGT	TTAGTAGCAA	CGTTTGTAAT	13560
ACCTTGATGAT	AGTGGTTCTA	CGGCTGCGAT	GAACCATGAA	CCGATAAGTG	AGAAGAGTTC	13620
CAAAGATACA	CCGGCAGTCA	AACCAAGAGA	CAACCATCCT	TTGATAACAA	GACGCCATTT	13680
ATCTGCATCT	GCAACACCTG	CAATTGGATG	TGGAGTTCCC	ATAATACCGA	TAACGATACC	13740
AAGGATGAAA	CCGATGAAGA	ATTTAGATCC	CCAGAAACCG	ATTTTCTTGT	TCAATTTAGC	13800
AGCATCAAAG	TCATATTTAT	CAAGGCCTGG	GAAGAATTTT	TCAAAAATCT	TATCCAAAAC	13860
CATGATAACT	GGTTCATCA	TGTAGTTCAT	GTGAGTTGAT	GTCATTGGTG	ATGAACTTGG	13920
GGCGTTAAGA	AGGTCATCAA	ATGTAGGTTT	CATCAAGTCA	GAGTTGATAA	TTTTCAACAC	13980
ACCGACAAGG	ACGATAGCTG	CTGTAGCAAT	AAAGAGTGAA	ACCCCTTGAC	TCACACCATT	14040
GTTATCAGCA	TACCATTTAA	TCAAGAGACC	TGTGATAGAC	AAGTGCCAGA	TATCAAAGAT	14100
ATCGACATCA	AGTGATCTG	TTTTCTTCAT	AGCTAGCATC	ACTATGTTGA	CAATCAACAT	14160

GATGAGCAAG AAGTATAGTG TCCAAGCAGA ACCCCAAGTG ATTGTAGCAA GTGGTGCCCA 14220
 ACCAACGTCG GTAATACTCA ATTGGATACC AGTGTTTTCA ACGAATTTG CTAGTGATGC 14280
 TGAGAAAGCA GTGTTTAGCA TACCGATGAT AGCACCGATA CCTGTAAGAG CGATGGCAAG 14340
 TTTGATACCA CCTTCAAGCG CTTTGGAGAA TTTCACTCCA AAAAGTAAAG CCAATACTGT 14400
 CAAAATGATT AACATGATGA CAGGTCCACC CATTCTAAG ATGGGATTGA AAACCTTTC 14460
 GATTAGGTCA AAGATTGCAT CCATAACAGT TCCTCCCTTT TTGATGTTAT ATGAATGTTA 14520
 ACAAATTAGA ATTAGCTTAA TCCGTGTTCT TTAATAGCTG CTTCAATATT GTCAAATACT 14580
 GGAGCGCTCA TTGCTGGGAT ACGGAATAAG ATTGGCCCAG CTTGATAAC TGGGATACCT 14640
 GGTCAAAAC CAAGGTCTGT TGCAGCGATT GGTGTAAAGA TATCGTAACC TTTCATAAGG 14700
 TCTTCGTTTA CATCTTTCAC CATGACTGCA TCACAGTGAA CATCATAACC ACGGTTTGAA 14760
 AGTTCTTCTT CTAGAGCACT TTTAATTTGG TGACTTGAGT TAACACCTGC ACCGCAGGCA 14820
 GCAAGAATTT TAATCATTTA GATTCCTCC GATTTATTT TTTAATAGAC AAGATTAAGC 14880
 GGTGCTTCA GCAATGTAAG TATAAAGGGC TTCTGGTTCA GAAATTTTG ATAGGTCTTC 14940
 AAGATGACCA TTCCTGTGA AGAAGTCCAT TAACTGAGCA AGAATGTTTCG TTTGACTTGA 15000
 ACTTGAATTA TTAATGATAA AGAAGAGTAG GGATACTTCT ACTTCCTTAT CAGGAGCTAT 15060
 CATATTGTGA AAAGTTATTG GTTTTCTAA TCGAACAACC ACCACTTCT CAGCTAGATT 15120
 ATGAACAATA TCTGTGTGAG GAATCGCTAC ATTTGGCAAG TCCTTTCCTA GAAATTCAT 15180
 ATCTAAACCA GTTGAAATG ACTTTTCAG CGTGATCAAG GCTTCACGAT AAGTTGGAGT 15240
 GACAATTTCT CGTTCTTCCA ATAAAGTTGC AACCTGATCA AAGAGTTGTT CTTGACTATC 15300
 CGCTTCTAAG CAAAACACAA GGTTTTGTG AAAGAAATAA TCTAATACCA TAAGTTTTTC 15360
 CGG 15363

(2) INFORMATION FOR SEQ ID NO: 140:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 28882 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 140:

TAAGACTAT TAATAGTGGA GTGAAATAG ATACGAACAA ATTGATTAG AAAATCAAAT 60
 GAATTTATAG AAATCTTTTA GCAGTTATGT TATCCTATTC TAGTTTCAA ACGCTATAGA 120

AGCAGCATTG	TGCTAGTCKA	GATTCAGTTT	ACTATACTAA	AACGAGTAGC	TTGAAATCAA	180
AAAACCCACC	CTCACAGGCA	GGTTTTATCT	GTATTATPCA	GCTAGATTAT	GCTTTACCTT	240
CTGAACCGAA	TACGTCGATA	CGTTCTTCAA	CCGATGCTTG	GATAGCTTTT	ACACCGTCAG	300
CCAAGAATTT	ACGTGGGTCG	AAGAGTTTTT	TCTTGTCGTA	TTCTGCTTCG	TTTGCTTCGT	360
AGTCACGAGC	AAATTTACGA	GTTGCGTTAG	CGAATGCGAT	TTGGCATTCT	GTGTAAACGT	420
TAACTTTGGC	AACACCAAGT	TTGATAGCTG	CTTGGATTTG	CTCATCAGGA	ATACCTGATC	480
CACCGTGCAA	TACGATTGGG	AATCCTGGAA	GAGCTTCTGT	CAATTTTTCG	AAGTGGTCAA	540
GGTCAAGACC	TTCCCAGTTT	ACTGGGTAAG	GACCGTGGAT	GTTACCGATA	CCAGCTGCCA	600
AGAAGTCGAT	ACCAGTTTCA	ACCATTGCTT	TAGCGTCTTC	GATTGGAGCC	AATTCACCTT	660
TACCGATGAT	TCCATCTTCT	TCACCACCGA	TAGTACCAAC	TTCAGCTTCT	ACTGAGATAC	720
CTTTAGCGTG	TGCTTTTTCA	ACAACCTCTT	TAGCCAATTT	AAGGTTTTCT	TCAACTGGAA	780
GGTGTGAACC	GTCAAACATG	ATTGAAGTAT	AACCAACTTC	GATACACTCA	AGTGCATCTT	840
CGTAGTGACC	GTGGTCAAGG	TGGATAGCTA	CTGGTACAGT	GATACCCATT	GATTCAACAA	900
GGTTAGCGAT	CAAGTTGCGA	GCAACTTTGT	AACCACCCAT	GTATTTAGCA	GCACCCATTG	960
AAGTTTGGAT	CAAAACTGGA	GCTTTTTTAG	CTTCTGCTGC	GCGCAAGATA	GCTTGAGTCC	1020
ACTCAAGGTT	GTTTGTGTTA	AATCCACCAA	CTGCATAACC	GTTGTCACGG	GCTGCTTGGA	1080
CAAATTTTTC	TGCTGAAACG	ATTGCCATTT	TATCAGGCCT	CCTGTATATT	TTTATGGGTC	1140
ATCCCATTTA	CATTGTTCAT	TTTATCACTT	TTTGCCAAAA	AAATCTAGTT	TTTCCCGCAG	1200
TTTCGATTGA	TTTTCTTCTA	ACTCCATCTA	TGTAAACCCT	TTCTCTCCCT	AGTCTTGGAC	1260
GACTTTTGGG	AAATCTATAA	AGAAGGTAA	ACTATCTCTC	TCCATCTCGA	AACGATAAGC	1320
TAATTTTTC	TGTTCTAATA	GACTCTTAAC	CACAAAGAGC	CCCATAACCAG	ACCCCTTGAC	1380
CTTGCGACTG	GCATTGTCAG	AAAAAGACTG	GGCTAGTTTT	TCTTGTTCCT	CTGAGCTACA	1440
GCTATTTTCG	ATAAAAAGTT	CTCCTTCTCT	TTCTCCAATT	CGAACTAAGC	CACCTGGAAC	1500
AGAGTGCTTA	ATGGCATTGC	TGATGAGATT	AGAAAGAAATC	AACTTCATAA	CTGATGGGTT	1560
TAGATAAGCC	TGCTGATGGG	TCAAACATTT	GTCTATCTGG	AGCTCTCTTT	CCTTGGCTAG	1620
CAAGGCATAA	TCTTTGACCA	GATTTTGCCT	CATCTGGAGG	AGGTCAATTG	TTTCCCTATC	1680
ATCTCGCAAT	TCCTGCACAG	AAGAGAGGGA	AAGTATCTGC	AGAACATGGT	GATTGAGTTC	1740
ATCCACAATC	CCCAAGGCAA	CTCCCAGATA	CTGGTCTCTA	TCCTTATAAC	GACCGATATT	1800
CTCTCTCATA	TTTTCGATTA	GGATTTTCAA	ACTAGCCAGC	GGTGTTTTCA	ATTCATGAGA	1860
AGCTCCTCGT	AGGAATTCGA	CCTTCATCTT	CTCCAGCTGG	AGAATGGCTT	CATTCTTTTC	1920

ATGCAAGTCC	GCAATAACAG	TCAAGAGATG	CTGGTAGAGG	CTATTGATTT	G TTCCTTGAG	1980
ATTACCTATC	TCATCCTTAG	AATCCACGCG	CAATCGCACT	TGGGAATCCA	GGTCCATCAT	2040
CCGACGGGTC	ACCCGCTTGA	TTTCCAAAAT	CGGTGCAACA	ATAGTCCGAG	CGTAGATGTA	2100
GGCCACCAAA	AGGGAAATCA	GAAAGGAGGC	CAGCAAGGTA	TAGGGAAGAA	ACTGGGAGACT	2160
GATTTGCTCC	GCTTCCTTTT	GTAATCCAT	GGAAGCTAGA	AACTGGAGAA	TCATAGTACC	2220
ACCGTCTTGC	GTTTTCACCT	CGCGCTCCTC	AATAAAGAGA	GAGGTTGTCT	GGCGGTCTGT	2280
GTCCAGAGGA	AGACTGTCCCT	TGACTTCTAA	CTTGTCTCTG	GTCATCTCAC	CTTTGACGGT	2340
CCCCTTGATA	TCACTAGTCT	GGGAATACAA	GTCTAACACT	TGCTCGATAC	TCTGCCTATC	2400
TTTCCCTTCT	AGGGACTGGG	CAATGGCTGT	TGCCTTTTGA	CCAATGGTTT	CCTGACGATG	2460
ACTCAGATAA	GTCGAAGGAA	AAAGAAAATA	AATAGCTAAA	TGAAGGCAGA	TAACCAGAAC	2520
ACTAAATATC	GAGAAGGTAT	AGATAAATAT	CTTTGCAAAT	AAACCTGTTC	GTTTCATTTT	2580
CGCTCCAATT	TATAACCAAC	ATTGCGCACA	GTGAGGATAC	AATCCAAGTC	TAGCTTTTTT	2640
CGCAATTCCT	TGATATAAAC	ATCAATAACA	CGGTCAAAGG	GAACCTCATC	TGTCGCTTTC	2700
CAGACGGCAT	CGATAATCTG	AGATCGAGTC	AAGGCCCGGC	CTTCATTTTT	CACTAGATAG	2760
TCCAGAATTT	CCAACCTCTT	GGCATTGATA	GGCACTTCTT	GACCTGCGAG	GCTTGCCTG	2820
TAGCTTTCAA	AGTCCACCTT	GGTATCCTTG	TAAGAAAAGA	TTCGTCTCTG	ATCGTAGTAG	2880
CGCTTGAAAA	TCGCGTCCAC	CCTCACTTTT	AAAAGGGAGA	GGGAGAAAGG	TTTTTCCAGA	2940
TAGCCATCTG	CAAAGAGGC	AAAGGCACTC	ATCTTGATAT	CCTCATCTTG	AAAAGCTGTC	3000
AACATCAAGA	CAGGAACCTG	ACTGGTTTTA	CGAATCTCAG	CTAGGACTTC	TAAGCCGTTG	3060
AGCTTGGGCA	TCTGGATATC	CAGTAAAACC	AGGGCCACCT	CATAGCTAGA	AAATTGCTCC	3120
AGAGCTTCCT	GACCGTCCGC	TGCCTCAATA	GTTTCATAGC	CACAATCCGT	CAAATAATCA	3180
CTGACCCCTT	CACGGATCAT	CTCTTCATCT	TCTACAATTA	AAATTTTCAT	ACTTTAACTG	3240
CTCTCTATTT	TTTATTTTTT	TTAGAATAAA	TACCTACCCT	ATTTTCTATT	ATAGTCTCTT	3300
GCTGGCCTTT	TGCTGCAAG	CAACTGACCA	CTAGATAAAA	CGTTGTGAAA	TTCTTTTCTC	3360
ATAAATTCCA	TAACCTTAGT	ATATTATATT	TAAGCACTAA	AGTACAAAGA	AAGCAACTGA	3420
AAGCAATGAT	TTTCACCACT	GCTTTCGGAT	TTATTTTGAA	TTGTAAAATA	GCCATTCCTA	3480
TCCACTATTC	TTGAATAGAA	ACACAAGATG	CAATCTTTAT	TCTAGACTCA	TTTTTTCAAA	3540
TTTATTCACC	ATCCAGCAAG	AGCTCTTTTG	GTTGTTTTCT	AAGGAGATTG	CTTGAAGCAA	3600
GCGCCATAAC	GAGAACCCTT	AGAACCAAGG	CAAGGACAAA	AATGATGATA	AAGTCTGATG	3660

TCTGAATGGA	AATGTCTAGG	CTCGACAAGG	TCTTGCTAAA	GCCATCTACT	TCTGCACCAC	3720
CACCAAGGTT	AGAGGCTTGA	GCCGCCTTAC	TAGCCTGTTT	GGCAACACCT	GAAGTCACAT	3780
TGGCAAGGAC	AGTGTTCCTA	ATTGCACGGG	CAGTGTAATT	AGCTAGGAAG	TAAGCAGAAA	3840
CTAGAGCAGG	GATAGCAATC	AAGATAGATT	CGGTGATGAA	TTGACCCAAG	ATACTTGCCCT	3900
GCTTGAGGCC	GATAGAGAGG	AGAATTCCCA	CTTCCTTGCG	ACGGGCGTTG	ATCCAAAGGC	3960
TGAGCAAGAG	GGCAAGGAGG	AGAACTGAGA	AGCTCAAGCT	ACCCAGAAG	AGGAGGTTGG	4020
CCATCTTGTA	CATACCAGAG	ATAGATTGCT	CAAGAGCTGG	GTAGTTAGAG	GAGCTCTTGA	4080
CGAGTGTGTA	GCTCTCCAG	TTGATACCAC	TGATGCCATT	CAACTCTTTC	ATAACATCAT	4140
CCAAGTCTT	GTCTGCTGTT	ACAAAGAAGG	TTGCGTCCCC	ATAAATGGCT	GTGTCTTCTG	4200
TGTATCCATA	AAGTTTTGCA	GCAGTGTGAA	TGTCTGTAAT	AGCTGTGTTT	TCGTAAAGTT	4260
CTTGTGAGTA	GGTACTGCT	GACTTATTAT	GACCATCAA	GAGTCCCTTG	ATTGTCACCT	4320
CAACTGTTTC	CTTGGCTCCT	TTTTCAATTAT	CTGCATCGTA	GATATTAGAG	TCCAGTTTAA	4380
CCTTGTCCCC	TACTTCCAG	CCGTGTTTGG	CTGCCAAGTC	CTTGTGCAAG	AGGATTTTAT	4440
CCTTGTGTC	GTTGGTTAAG	TGCTCTCCTT	CGACTAGTTT	ATAAGAACCA	GAGACAACT	4500
TGTCTTCTTT	AGAGGAGTCA	TTGACACCTG	TAATCATCAA	GCTACTTCCA	AAACGCTTGG	4560
CACGATCAGC	AGTGAGATTC	TTCTTGGTTT	CTGGCGTTTC	AATCAGGTCA	TATCCAGTCA	4620
AATCTCCGAT	AGCGTTGATA	CGTTTGACAT	AAGACTCAAT	GGCCTTGTTT	TCGGTGATTT	4680
TTTTGATGTC	TTCACCCCTG	ATATTCCCAG	CACCACGAGG	CGTTCCTTGG	TTGACGCGAC	4740
GATTGATTTG	CATGGAGAAG	CTATTGGTGA	TATTTTTAAA	GGTCTCCTGA	GAAGCCTTGG	4800
CAGTAGCTCC	CTTGATTGAC	AAGCCGACCA	AACTCAAGCT	CGCCATGAGG	AGAATAATCA	4860
GGAAGATGAC	AATCGATTTG	AAAAACTTCC	TTGTAACATA	GGCAAATGCG	TTGTGTAACA	4920
TAGATTC CCT	TTCTAGATTT	TGTTTTAATC	ATTCTATTAA	AATAAGCTCA	AATTATTTAC	4980
TAGTATTGCG	CGTTTCAGTC	AGTTTCTTAT	CCTTTAATTC	AAGTGAATA	TCTGACGCTT	5040
GTGCCACTTC	TTTACTGTGA	GTTACGACAA	TCACACATTT	ACCTGTTTTC	TGGGCAAGTG	5100
ATTTGAGTAG	TTGACAATA	TCTCCAGCAG	TTTTAGGATC	CAGATTTCCCT	GTTGGCTCAT	5160
CAGCTAGAAT	AACTGGAGCT	TCTGAGACCA	AACTGCGAGC	AATGGCAACA	CGTTGCTGTT	5220
GACCACCTGA	TAACTGGAGA	ACATTCGCT	TGATCTGGCT	TTCATCCAAA	CCAAGCTCAA	5280
GAAGTGATTT	CTTGCTTGCC	TTTTTGTGTA	CCAATCGGAT	ATTTTCCAGC	GGAGAAAGAT	5340
AATCTATCAA	GTTATAATTT	TGAAAGACCA	GGGAAATATG	GTGCATGCGA	TGGTAAGAAT	5400
AGCCCTTCTT	ACGAATATCC	TCTCCTTGAA	AAAGGATAGA	ACCTTCAACA	GGACTATCTA	5460

GACCAGCAAG TAGGGACAAG AGTGTGGATT TTCCTGCTCC TGA CTCCCCA ATAATACTGT 5520
 AAAATTTTCC GGGTTCAAAA TTATAATTGA TCTGATATAG GACTGCTTCA GCAGTATTCT 5580
 TATAACGGTA GGTAACATCT TGTAATTGTA ATAAAGTCAT GATTTCCTCT TCTTAACTAA 5640
 TAGATGATAA AATTTCTTTC GGTGATTTTC TAAATAAGAA TAGGAAACAA AGGGCTACAG 5700
 ATAAGCAACT AAGCAGA ACT AGAAAAACAT AGGATTCTGC AAAAGATAAG ATGCTAGTTG 5760
 ATAAACTGCT TGCTTTGGCT AGTGTATCTT GTAAGCTTGC CTGATCTCCA CTGCTAGTA 5820
 GAGTTTGGAG TAGGTAAGTT GTGATTGCGT TTCCTGCAAC AAATGCTGGA AGCAAAGCTC 5880
 CAAGAGATAC CAAA ACTACC TCTAAACAGA ATTGTAGGAA GATCGAGCTC TTGCCTTTTC 5940
 CAAGTGCAAG TAAAATCCCC ACTTCATAGA CCCGTTCTCT CAACCAGAGA GACAAAACCA 6000
 GAATTAAGGC TCCAGCTCCT GCTATCAACA TCCCATAAAG GAAGATGGTC AGGAAGGTTT 6060
 GGAAAGTTGC AACTGAGTCT TTGATTTGTT CAAAAGCCTT GTTTTCCTTT TCGACTTGGT 6120
 AGCCTTGATT TTCCAAGGCC AAGTTTTCTA CCTGCTTCAT GAGTCCGCTC ATTTCTTCTAG 6180
 GATTTTCTAC ATAGAAGCGT GCTGCAC TGA CTGAGCTTC ACTATTGCC AAAAGGGTTT 6240
 GGCTACTTTC ATAGTCTGTA AAGACTTGAT TTTCACTGAA GTCAGAAGAC AAGCCTGTGA 6300
 ATTTCTCTTG TTTTTTACCA GAAAAGATGC CGATAATCTC AACTCTACT GTTTGTCTT 6360
 TTCCAGATTC AGACTGACCA GCATCCAAGC CAATCTTGTC ATGAAGCGAA AGACCGTTCT 6420
 TCTTAGCCAA TTCTTCGTGG ATAAGGATTT TCTTGAATC CCCTTTTGA AGGTGTGCGC 6480
 CTCTTTTAG ATTGAAAGCC GAACTGGTAA AGGTTACATC CTGGATGAA TCCTCAAGAG 6540
 CCGTTAAGCT AACCAAGTTA TTGTCTGCAG CTGATAAATC ATCACGCTCC ACGCTCTGCT 6600
 CGCCAGTCAC TGCTTCTTG TCTTTTAGTT TTGCGACCGT CTCAAGTTCA GGAGAGACAT 6660
 TTCCAGCCC CTTAATCTTG CTACAGATG CTAGGCTGTA CAACTTGAAT GTCTGACCAT 6720
 TCTCTATCTT CTTAATAGAA AAAGATGTAT TGAGTGATTT ATAAAGATTG CTTTCTACTG 6780
 TTTTGTGGA CTTTCATCAGA GTCAAACAGG CTGAAATTC GGCCAATAAG ACCAATAAAA 6840
 TCAGAAATAA AATAAACTT CTCAGTCGCT TTCTGCTGAC ATAAGCCCAA GATCTTTGGA 6900
 TTGGATTCAT TTGTACCTC CATATTTGTA AGACTATTAT AAAACCCAAA TATGAAATAT 6960
 TTATGAAATA CGAAAAAAA ATATCGAGTA GGGGATAATC TCTAGCCCCT CTCACACCAC 7020
 CATACGTGCC GTTCGGCATA CGGCGGTTCA ACTA ACTTTT AACGCATGTC GTTCAAGGTA 7080
 ATAATCCAAA CACGAAACCA GTCCACGTTT TTCAAGGACT GGTTTGTATA TAGCACGTTT 7140
 AAGTACCGAC TTCTGAGCTA CTATAGTAGA TTGAAACTAG AATAGTACAC CTCTACTTCT 7200

AAAATATTGT	TAGAAATCGA	TTTGA CTGTC	CTGAACAATT	CGTCCTATTC	TTATTTTCATT	7260
TTACTATAAT	TGATAGTGGT	CGCCCCAGCC	AGATACCCTTA	TCTGCTATCC	ATTTAGGAAC	7320
CCCTAACTTA	AGCAATCCCC	ATAATCGTCT	CGATTTCTTC	TTCCATTGCT	TCCAGATAAT	7380
CACTCGTAGG	CGAGTACGCA	AGCGCTCATC	TATGCTAGTG	ACTATACTTT	TCATATTTAT	7440
AATTCATTC	TTTCGTTTCA	CTCAAGGCAC	AACACAGAAT	GAAAAAGTGT	TGTGATCTTT	7500
ATTTTGT TTT	ATAATAATAG	TGAGAAAACC	TATCACTACT	ACAAATCACG	GGGAGGTGAA	7560
TAAGTGAGTG	GTACAGCCAC	TACCTCGCAT	ATTTTGTCCAC	ATCATTTAAC	GGTACATAAT	7620
AAGTTGTACC	ATCTGAATAA	GTTGCTACAA	TATCATTTGC	ATGCTCTCCT	TCACCTTTAG	7680
CAAAGTTGG	AGCTCCTGCT	GGATGATTTT	TATTTGCCCTC	TTTCAATTTT	TCAATAATGG	7740
CATTTTCTCT	GTATCTTTTA	TATTATCAGG	ATTTTCTACT	AAGATTTTGT	CTGGATATGT	7800
CGGTTTAGCA	GAAACAATTT	TACTGT TAC	TTCTTTTTTA	TTCGAAGCAC	TTGTCCAGTT	7860
TCCAGCATT	TCTTTAGCAT	TAAATTTTAC	AGTAATTCCT	GAAC TAGGAA	CTTCAGTAGC	7920
AGGTTGATTA	TCAACATTAT	TCAACTTTAA	TTTCAAAAGA	GCTGTTGCAT	CAGACGTTTT	7980
ATCAATCGTT	ATATATAATG	ATGAAT TGT	ATTATAAACA	GTTCCCTCAT	ATTTAGCTGT	8040
TTGTGAGCTA	CTTGAAACAG	AACTGAAATT	ATACCCACTA	CCTCCCTGAT	TATCTTCAAT	8100
GCTTACGCTCT	AAATGAACTT	CCCCACTATT	ATTTGGCTTA	GCAACA ACTG	TTATAGTAAA	8160
ATAACATAAA	ATTTGCATAA	ATAGATTAGG	GAAATCAAAG	CAGCTTCTAG	GAATGTTTTA	8220
GCAGTCACAG	TGTACTTTCC	CAGCATCAAG	CCACTATAAC	TCTGCACATA	AAAATGGAGA	8280
AGATGGCAAT	CCTCTTCTCC	AAATATTAAC	TTCTTTTACAA	ACCAACTATA	GTTGACAAAG	8340
AACCTAAAAT	CAATGATAA	CACAAGGTCA	GGTCGGTCAA	CTCTTTCAAC	TGAAGCCCTG	8400
TCAACTCTTC	CCATTTATCA	ATCTTGATTT	GGAGAGAATT	GCGGTGCAGA	TAGAGTTGCT	8460
GGGCTGTTTT	AGTGAGAAC	GCACTATTTT	CCCAAAGAGA	GAGAATGATT	TCCTGAATCT	8520
GATCTTGATC	CAAAATCATC	TGGTGTAGAC	ATTCTTTGAT	TGGCTTCAAG	TCCACGAGTC	8580
TTTCTCCCAT	ACTCCAAAGA	TAGAGCTGAG	AAAAAGTATG	AACACCTTGG	TGACCCCTGAC	8640
GCCACCATGT	CTTGAACAAA	TCCCGCTCAG	CTTTGATTAA	GTCTGATAGG	GCTTGATGTC	8700
CCGTCTGAGA	CCAAACCTGA	CCCAACATGA	TAGAAAGACG	AAGTCCAAAG	TCATACTCAA	8760
CCGCTTCAAT	CGTATCACTT	AAAATATCTC	TTACAGAAGT	GTATTTGTCT	TGTTGAAGCA	8820
CGAAAACATA	ATCCTGAGCT	CCGACCTGTA	GCACTGTCTG	ACAATTCGGA	AAAAGAGTCC	8880
GCATCATATC	TAGCCAAGAA	GCCAGATTTT	CCTGCTGAAA	ATAAGAAAGA	TGGCAATAAA	8940
CCAACTGAAT	CTTTTAAAA	ACTTGCGGTG	CCTGTCCCTT	GCCCTCAACC	AGATAGGAAT	9000

945

ACCAAGGGTT	TAGCGAACGA	GCCTGCTCCT	GCTGGGTCAA	AAGGGCAACC	AACTGCTTTT	9060
CACGCTCGCT	GAGCCCAGCT	TCCTCCAGCA	AAATCCACTG	CTGAGAAGCT	AAAGGGAGCG	9120
TGAGATAGCC	CTCTTTCTCT	ACTGGTTGGT	CTGAAATCCG	AGCCTCAGGA	AACCAGTCTT	9180
GTAGTTCTTT	TGCCCTCATG	TTCTAGCCCT	CCACTTTTTG	GATGCACCAT	GAAACCAAAC	9240
TCTCAAGACG	TTCCAGATTC	TCAGTCATAT	GGAGATAGCC	CATAACCGCT	TCAAATCCCCG	9300
TGGACATACG	ATAAGTCACG	ACATCTGCAT	TTTTAGCCTT	TGTGTGGCTA	TTGGTATTGC	9360
GGCCACGTTT	GTAGATTTCT	TCTTCTTTTT	CCGTTAGGAC	CTGCTCCTCC	AACATGAGAG	9420
CAATCAGGCG	AGCCTGAGCC	TTGGCTGACA	CGTACTTAGT	TGCTTCTTGA	TGGAGTTTAT	9480
TGGGTTTGGT	CATACCTTTG	AGGATGAGGT	GACGGCGAAT	ATACATAGAA	TACACCGCAT	9540
CCCCCTCAA	GGCTAGCGCA	ATCCCGTTAA	TGAGATTGAC	ATCAATCACG	TGTCCACCTC	9600
ACTCCATCCT	TGGTATCAAG	GAGCTTAATT	CCTTGAGTAA	CCAATGGTC	ACGGATTTGG	9660
TCTGCTGTCG	CAAAGTCACG	ATTGGCACGC	GCCTCTTGGC	GTTTTTGAAT	CAAGTCTTCA	9720
ATCTCTGCAT	CCAAAAC TTC	CTCAACAAAG	ACAATTCCAA	AAATTTCTAA	CATATCTGCA	9780
AGAGCTTGCT	TGACACTTGC	ATCATAGTTC	CCTGAGTTGA	TCCATTTGGC	CATTTCAAAG	9840
ACAAC TGTGA	TACCGTTGGC	AGCATTA AAA	TCTTCATCCA	TAGCTGCTAC	AAACTTATCT	9900
TTAAAGTTTT	GTAAC TCTTG	GGCATCCACA	TTTCTGTAA	ATGGTTGTTT	GTAAGTATTC	9960
TTGAGATACT	TGAGATTGGT	CTCGGCATCG	CGAACTGCCT	TTTCCGTGAA	GTTGATAGGC	10020
TTACGGTAGT	GCTGGGTCGC	AAAGAAGAAA	CGAAGTACTT	GCCCATCAAG	AGTTTTAAGG	10080
GCATCGTGTA	CCGTAATGAA	GTTACCCAAG	GACTTAGACA	TTTTGACATT	GTCGATATTG	10140
ACAAAGCCAT	TGTGCATCCA	GTAGTTAGCA	AAAGCCTTGC	CTGTTTTAGC	TTCAGACTGG	10200
GCAATTT CAT	TGGTGTGGTG	TGGAAACTCT	AGGTCAGCTC	CACCACCGTG	GATATCAATG	10260
GTATCACCTA	AAATCTCTGT	CGACATGACT	GAACACTCAA	TATGCCAACC	CGGACGTCCA	10320
GGTCCCCAAG	GACTATCCCA	AGAAATCTCA	CCTGGTTTGG	AAGATTTCCA	TAGAGCAAAG	10380
TCTACAGGAT	TTTCTTACG	AGCCGTTTCT	TCATCGGTAC	GACCTGAAGC	ACCTAGCTCC	10440
AAATCTTCCA	AGGTTTTATP	AGCCAATTTA	GCATAGTTGT	GGGATTTTTT	TACACGGAAA	10500
TAGACATCCC	CTTGACTCTC	ATAGGCAAAG	CCTTCTCGA	TCAAGTCTTC	CACAAAACGG	10560
ATGATGTCTG	CCATAAACTC	CACTACACGC	GGATGGCGAG	TCGCAGGTTT	CACGCCCAAT	10620
GCCGTCACAT	CCTCACGAAA	GGCAGCGATG	TACTTATCCG	CAACCTCCTG	AGGCGTGATA	10680
CCTTCTTCCC	TGGCACGGTT	GATAATCTTA	TCATCCACAT	CTGTAAAATT	GGAAATATAG	10740

GCAACCTTAT ACCCACGGTA CTCAAAATAG CGACGAATCG TATCAAAAGC TACCGTCGAA 10800
 CGGGCGTTTC CTACGTGGAT ATAGTTGTAC ACCGTTGGCC CACAAACATA CATCTTGATC 10860
 TTGCCGTCCT CAATCGGGAC AAATTCCTCGC AAATCACGAG ACATGGTGTC ATAGATTTTA 10920
 ATCATAAATC ATAATCAGGA AAGCTGAAAT CCAAGAACAA TTAGTTTCAT CACTAAAAGT 10980
 TCAAGTAAAT TTCAGTCCGA ATATCTCTAC ACTTCGGAAT CCCTTGCTCC TTTCTCATTC 11040
 AGATAAACCA CCTGAGTCTG TTTGACAAAG CCAATTTTTT CATACAAACG TTTGGCACCT 11100
 ACATTGCTAT CTTCACCTGC AATCTGAAAT TCCTTGTCAT TTTGCTCAAT TAGTTGGTTG 11160
 ACGAGGGATT TTGCTAAGTA GCTTCCATAG CCTTTTCCAC GTTCAGGTTC CAATATTGCT 11220
 AAACCGTAGA GGTAAATTCGT ATTAGTCGAT AAATCAACCG TACAAGTTCC AATAACCTGA 11280
 CCAGCTTTTA ATAAAATATA TAGTCGGCTT TCTGGATCTT TCAGAGCTTC AGCGACATAT 11340
 CTATCCACAA CTCTCTCGA TTCATGTTCC TCTGAAAATG CCTGAAATTT TAATGACTA 11400
 ATTTGATCCT GATACGAAT ATCTGCTAAC AAAACTTCAA GATGGGAAAC ATTTGCTAAC 11460
 GGATAAGGTC TTCTATCCTT ACCTAACCAA GTTCTGTCTT CTTCATCCTC GATTAGTCCC 11520
 CAGTTACTGG CAAAGTCAGG ATGATTCTCT AAAAAAATAC GTTCTGTCTG AAAAGTGACT 11580
 GACCGAATGG GGAAGAAGC TGTTTCTCTC TCAAAACTAG TAAACAATGC ACGCGCAATC 11640
 CCCTGACGGC GATGACCTGG ATGAACCACT ATCGTCACTT CTACATCTTG GTCATCTGCA 11700
 TAGACAGTTA ATAAACCAAC AAGTTCGCCT TTTTCATAAT AAAGGAAAAA GCGGGCATG 11760
 TTTGGGTCAA AATTAAGCAT GTTAGAGAGA TAGGGATCGC GATAGGTACC GTCATAGTTT 11820
 TGGCAACAGT TAATTACTTT TTTGCCTCA GATAGCTCCT CTTGGCTTAA CTTGTTTCTT 11880
 GCTTGAATCA TATAGGTATC CTCTACAAAC CAGACGATCT GTGACTGGCA TCTTTAGCCT 11940
 GCTCGAGTTT ATTGACATAA TACTCTCGTT TTTCTTCGAC TTCGTGAATG ACAGGCTCAT 12000
 CTTTCTTACC ATGAAGACGG ACAATCTTGG CCGGAATACC GACAACCGTC ACGTCACTAG 12060
 GTACATCTGC TACGACAACT GCTGCAGCAC CGACCTTGGC ATTTTCACCA ATTTCCACAG 12120
 GCCCCATAAC TTGGGCATGG GCTGATATGA GGGCTCCCTT TCGTACAGTC GGATGGCGTT 12180
 TGCCACAGTC TTTCCCTGTT CCCCCGAGAG TCACTCCGTG ATAGAGAAGA ACGCCTTTTT 12240
 CAACAATCGC TGCTCTCCA ATCACCAGAC CAGAACCATG GTCAATAAAA ACACCTGAAT 12300
 CAATCTGGGC TCCTGGATGA ATCTCAATCT GAGTCCAAAA GCGCCAAAAC TGACTGTACA 12360
 TACGAGCTAA TAGTTTGAAG CCGTGCTTCC AGAGAAAATG CGAGAGACGG TGGGCCGCCA 12420
 AGGCCTTGAC ACCTGGATAA GTCAGCAAAA CCTCCAAAGT GGTGCGGGCC GCTGGATCAT 12480
 TTTCTTTTAC AATATCAATG GTTTCGCGCC ACCACCCCAT ACATTTCTCC TTTTCTTATT 12540

947

CTGAATCTTT	TGATGTTTCT	GTAATTCCTT	TCTTAGGTTT	GTAATCCTTT	TGATGACGTG	12600
GGCGGTGAGG	GCGCTCAGAC	TTTTCACCTT	TTTCATCATG	CTCAGGTTTT	GGCGGACGAG	12660
GTAGAAGAGC	CTTCATAGAG	GCATCGATAC	GGCCTTTTTC	ATCAATTTTG	ATAACCTTAA	12720
CATCAACTTC	ATCCCCGATT	TCTACCAAAT	CCTCTACACG	ATTGGTACGA	GTCCAAGCCA	12780
TCTCAGAGAT	ATGAACAAGG	GCATCTGTCT	TATCAAAGAG	GTTAACAAAG	GCACCAAATT	12840
TCTCGATACG	AACGACTTTA	GCACGGTAAA	CTTCATCCAC	TTTGGCTTCA	CGAACCAAAC	12900
CAGCAATAAT	TTCTTTGGCA	CGGTTAATAG	CATCTTGGTC	ACTAGAGTAG	ATAGACACAT	12960
TTCCTTCTTC	GTCATATCA	ATCTTAACAC	CTGTTTCAGC	GATAATCTTG	TCGATGGTTT	13020
CTCCACCCTT	ACCGATGACA	ATCTTAATCT	TGTCCACATC	AATCTTGATC	GTATCAATTT	13080
TCGGAGCAGT	TGGAGCCAAT	TCTGGACGAA	CTTCTGGAAT	GGTTGCTTCA	ATGACATCAA	13140
GGATTTCAAA	ACGCGCTTTC	TTGGCTTGAG	CAAGAGCCTC	CGTCAAGATT	TCTGCAGTAA	13200
TCCCTTGAAT	CTTGATATCC	ATTTGAAGGG	CTGTAATCCC	ATCACGAGTA	CCTGCAACCT	13260
TGAAGTCCAT	ATCTCCAAAG	TGATCTTCCA	AACCTTGGAT	ATCTGTCAAT	ACTGTGTAGT	13320
TATTTCCATC	TGAGATAAGC	CCCATAGCAA	TACCAGCTAC	TGGCGCCTTG	ATTGGCACAC	13380
CACCAGCCAT	AAGGGCAAGA	GTCCCGCAC	AGATAGAAGC	TTGAGATGAA	GAACCGTTTG	13440
ATTCAAAAAC	TTCTGCTACT	AGACGGATAG	CGTAGGGGAA	TTCTTCCAAG	CTTGGCAAGA	13500
CTTGAGCAAG	AGCACGCTCA	CCAAGGGCAC	CGTGACCGAT	TTCACGACGA	CCTGGCGCAC	13560
CGTAACGACC	TGTTTCCCCT	ACAGAATATT	GAGGGAAGTT	ATAGTGGTGC	ATAAAGCGTT	13620
TCTTGACTTC	TGGATCCAAA	CCATCAATGA	TTTGAGTTTC	TCCCATCGGA	GCCAAGGTCA	13680
AGACTGAAAG	AGCTTGAGTT	TGCCACGAG	TAAAGAGACC	TGAACCATGT	ACACGAGGAA	13740
GGAAGTCAAC	AACCGCATCC	AAAGGACGGA	TTTCATCGAC	CTTACGACCA	TCAGGACGCA	13800
CCTTGCTCTC	TGTAATTAAA	CGTCGCACTT	CTGCGTGTTC	CATTTGTTC	AAGATTTTCT	13860
CCACATCAGC	CATAATACGG	TCAAATTCCT	CGTGGTCCGC	ATATTTTTCT	TCGTAAACGG	13920
CAGTCACTTG	GTCTTTCACT	ACTTGAGTCG	CAGCTTCACG	GGCCAATTTT	TCTTCTACTT	13980
GAACTGCCTT	TTGGAGGTCA	CTGTTGTAGG	CTGCAATGAT	TTTCTTCTGC	AATTCAGCAT	14040
CCACGTGAAG	CAATTCCTTG	ACAGCTTCGT	GCCCTTTAAG	GAGCGCTTCC	AACATGATTT	14100
GGAAGGCAAT	CAATTCCTTG	ACAGCTTCGT	GCCCTTTAAG	GAGCGCTTCC	AACATGATTT	14160
CTTCTGACAA	TTCTTTGGCA	CCAGACTCTA	CCATGTTGAT	AGCGTGCTTG	GTCCAGCTA	14220
CTGTCAATTC	AAGAAGAGAT	TGCTCTGCTT	GTTCTTGACT	TGGGTTGATG	ATGATTTGGC	14280

CATCTACATA	TCCCACCTGT	ACCCCAGCAA	TTGGTCCGTC	AAATGGAATA	TCTGAAATAG	14340
ACAGTGCCAA	AGATGAACCA	AACATAGCAG	CCATTGGTGC	AGATGCATTT	TCATCATAAG	14400
AAAGCACTGT	ATTGATGACT	TGGACTTCAT	TACGGAAACC	TTCCGCAAAC	ATAGGACGAA	14460
TCGGACGGTC	AATCAAACGC	GCTGTCAAGG	TCGCATCTGT	TGAAGGACGT	CCTTCACGTT	14520
TCATAAAGCC	ACCAGGAAAC	TTCCCAGCCG	CATACATTTT	TTCTTCGTAG	TTGACTTGGA	14580
GTGGGAAGAA	ATCCCCAGTT	GCCATTTTCT	TAGACATAAC	GGCAGCAGTC	AAGACAGTTG	14640
ACTCACCGTA	ACGTACGACA	ACAGATCCAT	TTGCTTGCTT	AGCAACCTGA	CCAGTCTCTA	14700
CAATTAAGTC	ACGACCCGCA	AAAGTCGTTT	GAAACACTTG	TTTTGCCATT	TTAATCCCCT	14760
TTGGATTGAT	GAAATTATAC	GCCTTGCCTA	CAAAGATCAA	GATACCAAGG	ACGTCAAAAG	14820
CAAAGTAAAA	ATAGGAAACT	GACGAAGTCT	TCGATGAAGA	CAAGACAGTT	TATCTTTTTT	14880
ACACAGCTTT	TCGGCCGTGT	TCAATTACAC	AAGATATTTT	GGACGGTTCG	GCTTGCCGAA	14940
CATTTCTGTA	GAAAAATAGG	AAGGTGACGT	CGCACTCGAC	GAGTGCTAGG	AAGCTTATCT	15000
TTTTTCCTAA	GAAATGAGAC	CAAAATTCAA	GTCATCAAGA	TACCAAGCCG	TCAAGCAACT	15060
CAAAGGAAGA	TAGGAAATCG	AACGACGGAG	CGACTACTCC	TAGGGAGATT	TATCTTTTTTC	15120
CACAGAGTTG	TAGGCAAGTT	CAGTTTTCAA	GATACATCAT	TAGAAAGGTT	TAATACTAAA	15180
GTATCTAAAG	CTTTCACGCT	AATCGCTATC	GGGCGATTAG	CTAAATGCTT	TACTAACTCT	15240
CTCGTCAAT	AACATCGATT	TGACTCACTC	GTGTCGTAA	ATCTTACAGT	TTAAATGCAT	15300
TGTATTATTT	AATACCTTCA	TCTTTGTATC	AAGTACGTAC	AGAATTTATT	TTATCATATT	15360
TTTCTTAAAA	AGTGAGGTCT	TTACCATTAA	AAAGGAACCA	TTCCCCTCAC	CTGAGAAGAA	15420
TGGTTTGCTT	TTATTATCCT	AGAGACTGGT	GATTAAACAA	GGCATGGGTT	GCTTGATGGA	15480
TGTATTTTGC	TGTATCAGCA	TTATTATCG	TATAGAGATG	CACACCGGCA	ACATCCTGAG	15540
TTACCAAGTC	CACGATTGG	TCCACTGCAT	AGGCAAGTCC	TGCTGCTCTG	AGCGACTCAG	15600
GGTCATGCTC	ATACTTGCT	AAGATGGCTT	TAAATTTGCG	TGGAAGATGG	ATATTCTCAC	15660
AAGTCTTCAA	GAGTCGGAGA	GCCTGATTT	GATTCAGAAT	TGGCATAATT	CCTGCATGAA	15720
TGGGAACATC	AATCCCAGCC	AAGATACACT	TGTCCTGAAA	ATCATAGAAG	CGCTCATTTGT	15780
CAAAGAAGAG	CTGAGTTACG	AGGCTCGAAC	AGCCTGCATC	CACTTTCTTC	TTAAGATTTT	15840
GAATATCTGA	AATCTGATTT	GGCGAATCTG	GATGCCCTTC	TGGATAGCAA	GCTCCAATAA	15900
TATCAAAGTG	AGGGGTTTGT	TCCTTGATAA	ACTCAATCAA	GTCGGTTGCA	TAGCGGAAAT	15960
CCTTTTGTGG	TTCCACGTCT	GGAATAATAT	CCCCACGAAG	AGCCAAGATT	TTCTGCACCC	16020
CAACTTTGTC	CAAGTCAGCA	ATAGTTTCAG	CAACCTTGTC	CTTAGTTAGA	TAAATAGCTG	16080

GCAAGTGGGC AATGGTCGGA ATCGCCAAAT CATTTTGGAT AAAGTCAGCC AAACGAACCG 16140
TCGTTTCCTT GATATTAAAT TTATTATTGC TGGCAGTTAC ACTGATAAAA TGGGGAGCCA 16200
ACTCCTGCAT ATCCTGCAAG GCTGAAATAA TGTATCATT ACCCACGGCT GGGTTTGGAG 16260
GGAACACTTC AAATGAGAGT GACGGTGTFT GGCCTGACAT ATGTAATAAC CTTTCTAGT 16320
TGATTTCTTT TTGAACAACC ACTGTATGGA GAGAAATCCA ATCTTACAAT TTCTCACGGC 16380
CAGCTTTAGC TGCTTCAACA AGGCGGATCA AGCTTTCCTT TGTTCCTGGG ATACCACGTG 16440
TTTTCAAACC ACAGTCAGGG TTGATCCAAA CTTTCTTGCT TGGCACTTA GCAAGGATGG 16500
CTTCGATTGT GTTGTGATT TCGCCTTCAT TTGGTACACG AGGTGAGTGG ATATCGTAAA 16560
CCCCAGGTCC CACTTCTGTT TGAAGTTT TCGCTTGGAG TTCGTCCAAG ATTTCAAGGT 16620
TTGAACGGTT AGCTTCAAAG GAAATAACGT CTGCATCCAT GTTATCGATA GCTGGGATGA 16680
TATCTGTAAA TTCTGAGTAA CACATGTGAG TGTGGATTG TGTGTCTGGC GCTACTGTTG 16740
AGTGTACCAA GCGGAAGGCA GGAATAGCCC AGTCAAGGTA GTCTTCGTAC CAGTCGCTAC 16800
GGCGGAGTGG CAATTTTCA CGAAGAGCAG CCTCGTCGAT TTGGATGATT TTCACACCAG 16860
CAGCTTCAAG GTCAAGTACT TCATCCTTGA TAGCAAGGGC GATTTGGAGA GTTGAATCCT 16920
TGATAGAGAT GTCTTCACGT GGAATGACC AGTTAAGGAT GGTAACAGGT CCAGTCAACA 16980
TACCTTTAAC AGGTTTGTFT GTACGACTTT GTGCATAGCT AGACCATTA ACAGTGATAG 17040
GGTTAAGACG AGTGACATCA CCCAGATGA TTGGTGGTTT TACCCACGC ATACCGTATG 17100
ATTGTACCCA TCCATTTTGA GAGAAGAGGT ATCCTGACAA GTTTTGACCG AAGTACTCAA 17160
CCATGTCATT ACGCTCAAAT TCACCGTGA CAAGGACATC AAAGTCAATA TCTTCTTGCC 17220
ACTTGATCCA TFCGTCAATC GTTTCAGCAA GGAAAGCGTC GTACTCTTTT TGAGACAATT 17280
CACCTTTACG GTAAGCCAAA CGTTTGGCAC GAACCTCTTT TGTGAGGG AATGAACCAA 17340
TCGTTGTTGT TGAAGAGCT GGAAGTTTGA AAGCTTCTT TGGATAGCT TCACGTTCTG 17400
CAAAGGCTGG CAAACGAGTG TAGTCTGCGT CTGTCAAGCC AGCGATACGC GCACGAAGTT 17460
CAGCATTTTC ACCAACACGC TCAGTCGCAA AGAGTTCTTT GTTGGCTGCA AGAGCTTCTG 17520
AACCTTGACC ATTTCCGATA GCATCCAAGT CACGGATTTC ATCCAATTT TCAACTGCAA 17580
AGGCAAAGTG GTTCAAGAGT GCTGGTCAA ATTCTTCATT AGCAGTTGTA AATGGCACAT 17640
GAAGAAGTGA GCAAGAGCTT GTCAAAACGA TGTTTTCAGC TGAATTTGC TCAAGAACAG 17700
CCAAGCTCTT TTCGTAGTTG TTGCGCCAGA TGTTTTTACC ATTGACAATA CCTACATAGA 17760
GAGTCTTGTC AGCTGGGAAG CCACCTTAA CGAGTTCAAG AGTTTTCTTA CCTTCAACAA 17820

AGTCAAGACC	GATAGCATCT	ACTGGTAAGT	TTACAAGGTC	AGCGTATACG	TCACGAACAT	17880
CACCGAAATA	AGTTTGAAGC	AAGACTTCAA	GACCTTTTTT	GTCAGCCAAG	AGTTTGTGT	17940
AAAGGTTCOA	GAAGAGAGCT	TTTTCTTCAG	CTGTCAAGTC	TTTTACAAGA	GCCGCTTCAT	18000
CCAATTGGAT	GCGAGTCGCA	CCAAGTTCAG	CCAATTTAGC	AAAAACTTCT	TGTAAGCAG	18060
CCACTAAGCT	ATCTACGAAG	TCGTCTGCTT	TCACGCCTTC	TTCAAAGTCT	GACAATTGAA	18120
GGAAAGTGAA	GGGACCTACA	AGAACAGGAC	GAGTGTTCOA	TCCAAGTCT	TTGGCTTCTT	18180
GGAACTCATC	GAAAATCTTG	TGACCAGCCA	ATTTTACTTG	AGTGTCTTTT	TCAAATTTAG	18240
GAACGATGTA	GTGGTAGTTA	GTGTTGAACC	ATTTCTTCAT	TGGAAGGGCG	CGAACGTCCC	18300
CTTTTTCTCC	CTGGTAACCA	CGTCCCAAAG	CGAAGTAGCG	CTCAAGGTCA	GACAAGTCCA	18360
AGTTTTGAAC	GGATGCAGGC	ACCACGTTGA	AAAGGAAAGC	CGCATCTAGG	AAGTTATCAT	18420
AGTGAGAAAA	GTCATTTGAT	GGAATTTGAG	TGATGCCTTT	TTCTTTGACA	ATGTTCCAGT	18480
GTTTAGCAGC	CAAGTCTTTT	GCTGCTGCTA	AAAGTTCCTC	TTCTGAGATT	TCTTTTCTAA	18540
AGTATTTTTC	AGTTGTAAAT	TTTAATTCAC	GGAATTCGCC	CAAACGAGGG	AAACCGATGA	18600
TTGTAGTTGA	CATGATGTGT	CCTCCAAAAT	TTGTTGTTGA	AACTATCTTA	ACAGAAAAGA	18660
AAGCGTCTGT	ATAATTGTAA	AAAATTAGGG	TTTGATATAG	TTTGAAACTA	TATATCTGTT	18720
TCGGACAAAA	GAAAAAGACT	TGAAGCAAAC	GTCTCAAATC	CTTTGTAATT	CTFACTTTAC	18780
AGCTATATTC	CAATTAGAAT	ACTAAAACAT	GTTATTAGTA	ATTCTTATAA	GTGACTATGA	18840
CCTGTTATTA	GAAAAGACTA	TAAGTGAATC	TAGTCAACTT	TTTCCCTGTT	CAAGTGGGAC	18900
GATTGCTAGT	GTCTTTCTTA	AACTGGCTAG	GACTTTTAAAG	ACTGTATCCA	ACTGAGGACT	18960
AGTCTTTCTT	GTCTCCATCC	TAGCTATGAC	AGGCTGGCTT	ATTCCACTGA	CTTCTTCCAG	19020
CTTTTTCTGA	CTGATTCCTT	GTTTATACCT	AGCCTCAATC	AACTCGCTCA	TGATAGCCAC	19080
TCGCATATCA	CTTTCAAGGA	TTTCCCTCCTT	GCTAAAGAGC	TCAGATGGAC	ATCCTTCCAA	19140
TTACTCCCAA	TAGCACTATT	CTTCATCACT	TAACCCTCCTT	TTTTTTACGT	CTATGTATTT	19200
TTAAAAAAT	GAGCGAATTA	TGATTCGATA	GATTGACCAG	TGGGTTTAAA	GTTGGTGCTA	19260
GCCTATTTCT	TAAGCGATTT	TCCTTTTCTA	GGATAAAGCA	GTTCCCTGCTT	GCTTAACCCC	19320
AATTTTCCAC	GATGAATCCA	ATAGTAAATG	GTTGAAATTC	CCACGTTAAC	CCCTTTAGCC	19380
ATCACCATCA	TTTCAGGCGA	AAATTTTGG	TTATGTTTTT	GGTTATGTAT	AGTGGAGAAT	19440
CTTTTCTTTT	AGTTTCTTAA	GACTGTTGAG	CGTAGTCGGC	AGAATAAATC	TCTTTGAAGC	19500
GCCCTTTTCC	AAGACATTGT	CGGACTGTCC	CACGCTTGAT	TTCAGTGTGG	ATAGTTTGAG	19560
GAGCTTTTCC	AAGTAGAGAG	GCAATTTCTC	TATTTGATTT	TCCTTCTTTT	TTCCATCGTT	19620

951

CGATTAAGCG	ACGGCTATCG	ATTGTCAAAT	GTTTGCCTTT	TGTAGTATAA	TTGTCTTGCA	19680
TTTCTGTGCC	TTTAAATCAT	TTCAATCTTA	AATTGGACTT	TTTTTACTTG	GGTTGTA	19740
AATCTATGAG	GAAGACAAGA	AAAAGAATAT	CAATCAAGTA	AAGTCACAAA	GTCACATTAG	19800
CTCCGAGCAA	CCATTGCAAA	TTGAGGTA	CACACAATGA	TTAAAACATT	TCTCTCTGCC	19860
CTTTCGGTCA	TTCTCTTTTC	TATCCCTATC	ATAACTTATT	CTTTTTTCCC	ATCTTCTAAT	19920
CTTAACATTT	GGCTATCTAC	CCAACCTATC	TTGGCACAGA	TTTATGCCTT	CCCCTTAGCT	19980
ACTGCAACTA	TGGCTGTAT	TTAAGTTTC	TTATTTTTTT	TCCTATCTTT	TTACAAGAAA	20040
AATAAACAAA	TACGGTTTTA	CTCTGGCATT	TTGCTCTTAC	TATCGCTCAT	ATTACTATTA	20100
TTCCGGAACAG	ATAAAACCCT	TTCTTCTGCA	TCAAATAAGA	CTAAAACCTT	AAAATTAGTA	20160
ACTTGGAACG	TCGCTAATCA	AATAGAAGCA	CAACATATG	AGCGAATTTT	TAGCCATTTT	20220
GACGCCGATA	TGGCTATATT	CCCTGAACTA	GCTACCAATA	TCAGAGGTGA	GCAAGAAAAC	20280
CAGAGAAATCA	AACTATTGTT	TCATCAAGTT	GGACTTTCTA	TGGCCAACTA	TGATATTTTC	20340
ACTTCTCCAC	CTACCAATAG	TGGAATAGCT	CCTGTGACTG	TGATTGTCAA	GAAAAGTTAT	20400
GGTTTCTATA	CAGAAGCTAA	AACTTTTCAT	ACAACACGGT	TCGGGACAAT	TGTATTACAT	20460
TCGAGAAAAA	AAAATATACC	AGATATCATT	GCCTTGCCATA	CTGCGCCTCC	TCTGCCAGGT	20520
TTAATGGAAA	TCTGGAAGCA	AGACTTAAAC	ATCATTCATA	ATCAATTGGC	TTCAAAATAT	20580
CCAAAGGCTA	TTATTGCAGG	TGATTTTAAT	GCAACTATGC	GTCATGGAGC	ACTTGCAAAA	20640
ATAAGCTCTC	ATAGGGACGC	ATTAAATGCA	CTGCCACCTT	TTGAAAGAGG	AACTTGGAAT	20700
AGCCAAAGTC	CAAAACTTTT	TAATGCAACA	ATAGATCATA	TTTTATTGCC	TAAAACCAC	20760
TACTATGTTA	AAGATTAGA	CATTGTAAGT	TTCAAAACT	CTGATCATAG	ATGTATTTT	20820
ACAGAAATCA	CATTTTAATT	ATTTTATATA	AAATCACCCC	TCTAATGTTC	ATAAACTAGA	20880
GGGGGAATTT	GTATCCTACT	ATCGTTTAAC	GCACTTCTGC	ATTGACTTTT	TCTTCGAGAG	20940
ACGCTTGAT	TTTTTCCATA	TAGCGTCCGA	CTTCTTCGTC	CGTTAAGCTG	TCTTCTGGAT	21000
TTTGGAAAGT	CAAGCTATAA	GCCATTGACT	TCATACCAAG	TCCCAGTTT	TCACCTGAGA	21060
AGACGTCAAA	GAGTTTGATA	TCTGTCAAAC	GTTTCACGCC	GGCAGCTTGG	ATAGCATCTA	21120
CAACTTCTTG	GTGAGTCACT	TCTGCCTTGA	GGAGAAGGGC	AACGTCACGG	CTGACTGCTG	21180
GGAATTTGGT	GATTTCCACA	AATGGAACAG	CAGGTTGGAG	CGCCCCTTCG	ATGGCTGAAA	21240
GGTTAAGCTC	AGCTACATAC	GTTTCTGGAA	TATCGTAAGC	CTTGGCAGTG	ACTGGATGCA	21300
CTTGGCCAAG	GAAACCAAGA	ACTTGGTCAC	CGAGTGAAAT	CACGGCTGTA	CGACCTGGAT	21360

952

GAAGGCTAAC	GATTTCAGAT	GTTGCTGTAT	AGGTTACTTG	GAGTCCCAA	CGAGTAAATA	21420
GGGCTTCAAG	GATTCCTTA	GCATAGAAGA	AATCAACTGG	AACTGCTGCT	GTTTGGAAAT	21480
CTTTTTTCAGC	AACCAAGCCT	GTCAAGGCAA	AGGCAAAGCT	GTTGATCTCA	TTTGGAAAGTT	21540
CTTCTTTTGG	ATTACCTGTT	TGTTCAAAGA	CTTTTCCAAT	CTCATAAAGG	GCCAAGTTTTT	21600
TATCTTACG	AGCCACGTTG	TAGGCAACGG	TATCAAGGAT	CCCTGAAATC	ATATTTTGAC	21660
GGAGGACTGA	ACGATCCACA	GTCATTGGCC	ACATGAGTTC	AGTAAGGTTA	CTTGTTGAG	21720
CTGTGAACTC	AACTGCTTTT	TCAGGAGTTG	TCAGAGCATA	GGTGATGAT	TCTGTCAAAC	21780
CTGCTCCTTC	AGCAATGGTA	CGAACTTGAC	GGCGGAGTTT	TTGTATCACA	GTCAATTCAC	21840
CAGCTGTACC	ATCGTCTTTT	GGAAGGCTGG	TTGGCAAGCG	GTCATATCCA	TAGATACGAG	21900
CGATTTCTTC	AAAGAGATCA	GCTTCGATTG	TGATATCCCA	ACGACGACGT	GGTACGCTGA	21960
CTGTAAAGCT	GTCTGCATTT	CCAGAAAGAC	CAAAGCCAAG	ACGACGGAAG	ACGTCTTCTA	22020
CATCAGCATA	AGACAGCTCA	GTTCCGAGGA	CACGGTTAAC	ATCAGCAAGG	GTTGAAGAAA	22080
CTTCCACATC	AGAGGTATCA	AGCTCACCCG	CTGAAACGAT	ACCCTTACGC	ACCGTCGCGC	22140
CTGCAAGCTC	TGCAATCATG	CTAGCTGCCG	CATCAAGGGC	TTCATTAACT	GTTGCCACAT	22200
TAATTCCTTT	TTCAAAGCGA	GAAGATGACT	CAGAACGAAG	G TTCAGGCGA	CCACTTGTCT	22260
TACGGATAGA	TTTGCCATTA	AAAACAGCAG	CTTCAAGGAT	AACACGACTA	GATTTTTTCAG	22320
AAATTTCTGT	AGCCTGACCA	CCCATAACAC	CGGCAAGGGC	TACTGGTTTG	TCAGCAACTG	22380
TAATCACGAG	GTCTGTCTCA	GCCAAGTCTC	GTTCTTCACC	GTCCAGGGTC	ACTAATTTTT	22440
CACCATCACG	CGCTTCACGC	ACACGGATGT	CAGTCCCTTC	AAATGTGTCC	AAGTCAAAAG	22500
CATGCATAGG	TTGACCAAAG	TAGAGCAGGA	TGTAGTTTGT	CACGTCTACA	ACGTTATTGA	22560
TGGGACGGAT	GCCTTCGTTT	ATGAGAAGGT	TTTGCAACCA	TTGTGGACTT	GGTGCGATAG	22620
TCACATTGTC	CAAGATACGA	GCTGCATAGT	AAGGCGCCTT	GTCTGTCTCA	ATGCTGACAG	22680
AAAGGGCATC	TGCCCGAGCT	TCATTAGTTT	CTGTTAGAGT	AAATTTTTTA	AAGTTGACTG	22740
CCTTGTCATA	GATGGCTGCC	ACTTCGTGAG	CCACTCCACA	CATAGAAAGG	GCATCTGCAC	22800
GGTTTGGTGT	GATGGAAAGT	TCGATGATTT	CATCATCCAA	GTCTAGGTAA	GAAAAGACTT	22860
CCTCACCTGG	CACGGCATCT	TCAGGCAAGA	TTTGATGCC	ATCTGCGAAT	TCCTTAGGCA	22920
CAACTGAGTC	AGAAATTCCC	AATTCACCAA	GTGAACAGAT	CATTCCAAGT	GACTCCAAAC	22980
CACGGATTTT	TCCTTTTTTG	ATTTGTAGT	TATCAGCGAT	ACGAGCTCCT	GGAAGAGCCA	23040
CCATGACCTT	GATCCCAGCA	CGCACATTTG	GGGCACCACA	AACGATCTGA	CGCTCTTCTT	23100
CTTCGCCAAC	GTTAATCTGA	CAAACATGGA	GGTGAGTCTC	TGGCACATCT	TCGCAAGACA	23160

953

AGACCTCACC	GACGACAATT	TTTGAGAGAC	CAGCAGCTGG	TGATTCGACA	CCCTCTACCT	23220
CGATCCCTGT	AGTTGACATT	TTTTCAGCCA	ACTCTTGTGA	TGGCACATCA	ATGTCACCA	23280
ATTCFTTTAA	CCATTTATAA	GATACAAGCA	TAATTTAGTT	CTCCAGAATG	ACAGTTGTCA	23340
CTCTAGTTCT	TTTCCTTTCC	TATCATTTCA	ATAGAAGAAT	CCTCTTCTTA	CCTTAATTTT	23400
TTTCTCAGTA	ACCAATCCGT	ATCTACTTTT	TGACCAACCA	TAAAATGATG	TTGGCTAAAT	23460
TTTTCAAAAC	CATATCGGTT	ATAAAACGCT	TGAGCTTTTG	TATTATGCTC	CCAAACACCT	23520
AGCCAAGCCC	AAGAAAACCT	ATTTTTTGTA	GCAAGTTCOA	GTGCGAATTC	AAACAGTTGC	23580
TTACCTAGTC	CAAATCCTTG	GAATTTTTGT	AGCACATAGA	GACGTTGAAT	TTCAAAAGCG	23640
TCCTCTAATF	CTCTCTCAGT	TTGAGCACTT	CCCCAGTTGA	CTTTGAGAAA	ACCAGCTATC	23700
TCCTCCTCAT	GCATAATGAA	ATAGGTTTCA	GAGTCAGGAT	TTCCCAACTC	AGTTGACAAA	23760
GTTTTCAGAC	TATAAGCCTC	TTCAAAGTAT	TCCTGTAACF	GCTCTTCCGT	ATTATCATA	23820
GCAAAGGTTT	CACGAAAGGT	TTGTTTGGCA	ATTTTAGCCA	ACACCTCAAC	ATCTGCCATT	23880
TCTACTTTTC	TAATCATTAT	TTAAACTGTT	CTGAGAAGCG	GACATCTCCT	TGGTAGAATC	23940
CACGGATATC	GTTGATTCCA	TAACGGAGCA	TAGCTACACG	CTCTTGTCOA	AGACCAAAGG	24000
CAAAGCCAGA	GTATACAGTC	GCATCGATAC	CACTCATTTT	AAGGACACGT	GGGTGAACCA	24060
TACCGGCCCC	CATAATTTTC	ATCCAACCTG	TTTTCTTACA	TACATTACAG	CCTTCTCCAC	24120
CACACTTGAA	GCAAGAAACA	TCCACCTCAA	CAGATGGCTC	TGTGAATGGG	AAGTAAGATG	24180
GACGCAAACG	AATTTGACGC	TCTTCACCAA	ACATTTTTTG	GACAATCAAC	TGAAGCGTTC	24240
CTTGAAGATC	AGCCATAGAG	ATATTTTTCC	CAACTACCAA	GCCTTCGATT	TGGTGGAATT	24300
GGTGACTGTG	GGTCGCATCG	TCCGTATCGC	GACGGAAGAC	ACGCCCTGGC	GAGATCATCT	24360
TCAAAGGACC	TTTAGAAAAA	TCATGGGCAT	CCATAGCACG	CGCCTGAACF	GGAGACGTGT	24420
GGGTACGGAG	CAAGATTTCT	TCAGTGATAT	AGAAAAGTAT	CTGCATATCA	CGAGCTGGGT	24480
GGTCTTTTGG	AAGGTTTATA	CGTTCAAAAGT	TATAGTAGTC	TTGCTCCACT	TCAAAAACCAT	24540
CCACGACTTG	ATAACCCATA	CCGATGAAGA	TATCTTCGAT	TTCTTCACTG	GTTTGTGTCA	24600
AAACGTGACG	GTGACCAGTC	GCAACTGGAC	GACCTGGAAG	CGTCACATCT	ATACTCTCGC	24660
TAGCCAGTTG	AGCCGCGACT	TTCTTTTCTT	CCAAGAGCTT	AGCTGTTTCT	TCAAAAGCAG	24720
CAGTCAAGAC	ATCACGAGCT	TCATTGACGT	GTTTCCCAGT	GATTGGACGC	ATCTCAGCAG	24780
AAACATCTTT	CATCCCTTTG	AGGATTTTCA	TGAGCGAACC	CTTTTTACCA	AGGACAGAGA	24840
CACGCAAATC	TTGCATCTCT	TTTTCATTTT	CAGCAGTAAT	CTGCTTCAAG	CTAGCCAGCG	24900

TTTCTTCGCG	AAGCGCTTTT	AATTGTTCTT	CAATAGTTGA	CATATTTCCCT	CCATCAGTCT	24960
CTCGTAGATA	AAAAGAAAAC	CACATGCCAA	AAACTCCACT	CGGAGCGTTG	ACACGCGGTA	25020
CCATCCGTTT	TCATCTGACA	AGTCAGACCT	TCATTTCTAA	ATCCATGCCG	AAGTGAATTC	25080
ACCCAGCTTT	CATATAGAGA	GCTTGCAGTC	ACGGCTCTCC	TCCCTGATAT	ACTTCCCTTG	25140
GGCTACTAGT	CTTTCAGATT	CCTATTCAAT	TACTACTTAG	TTTATCAGAT	TTTTACCATT	25200
CTTGCAAGAC	CTATCTTACT	TCTGCTTGTT	AGCTTATTCT	TATCTAAATT	TATATAAACc	25260
TTATCTAAAT	TAACTATTTA	TAATTTTGT	AACAAAATTA	AATTAATTGA	CACTCCCCTA	25320
TAAAATAAAG	AAGTTTAGAA	TTTAATGTCT	TCCAAACTTC	TTTATTCCAT	ATTTAATGAA	25380
ATGCCACCTT	AACCGTGATA	ATAGCTAGTC	ATCAATAAAA	AACTATTTGA	ATAAGGATTC	25440
TCCATTTGAT	TCAATCACTT	CTTATAACCA	AGTAAAAGAC	ATTTTCTTAT	ATCGATTTAA	25500
TGTACCACTT	CCATCATCGT	TTCGATCAAC	ATAAATGAGA	CCGTACCTTT	TAGAAAGTTG	25560
TGCAGTGGAC	ATAGAAACAC	AGTCAATACA	TCCCCAAGAC	GTATAGCCCA	TAATTTCAAC	25620
ACCATCCTGT	AGAGCTTCAG	CAACTTGCAA	TAAATGTTCT	TTCATATACT	GAATTCCTATA	25680
ATCATCTTGG	ACGGTTAAGT	TATTAAGTTC	ATCTTTTATT	AGTTGATCTT	TAGCACCTAA	25740
TCCATTTTCT	ACTATAAATA	ATGGGATTTG	ATAACGGTCA	TAATATCTAT	TTAAAATTAT	25800
ACGTAGTCCA	ATGGGATCAA	TTTGCCATCC	CCACTCTGAA	GACTCTAAAT	AAGGATTTAC	25860
TAAACCACCA	ATAATATTCC	CTTCTCCTGA	ATTATACTGT	GTTGGAAGAG	CAGATTGAGT	25920
CACACTCATG	TAATAGCTAA	AGGATAAAAA	ATCTACGGTA	TAATTTTTTA	ATAACTCTGC	25980
ATCTTCAGCT	GCAAACCTCA	TGTTAATGTC	ATTTTCCTTA	AAATATCTTT	TTGCATAAAT	26040
CGGATAATAA	CCTCTAACAT	GCACATCTGA	AAATAGATAA	TTTAGATTCT	CATACTCATG	26100
AGTCGCCCAT	ACATCTTTTG	GATTTGGAGT	CATTGGATAA	GCTGGCATAG	CTAATACCAT	26160
ACATCCCACC	TTAAACTCTG	AATTAATCTC	ACGAGCAATT	TTTGTAACCA	AACTTGAGGC	26220
GACTAATTCA	TGATGTATAG	CTTGATATAA	TTCTTGTTTC	GAAAGATTCT	CCTTAGGTAT	26280
ATCTATTCCT	CCACTAGTAA	ATGGTAATTC	CAAAACAGAG	TTTACTTCGT	TAAATGTAAG	26340
CCAATATTTA	ACTTTATCTT	TATACCTTTC	TAAAACGTGT	CGAGCAAATT	TTTCATAAAA	26400
ATGAATCATT	CTCCTATCAA	CCCATCCATG	ATATTTTCTT	GCTAAATATA	ATGGAGTCTC	26460
ATAGTGTGAA	AGAGTTACAA	GTGGTTCTAT	CCCGTGAGCA	TGTAGTTCAT	CAAACAATTC	26520
ATCATAATAT	TTCAACCCAG	CTTCGTTAGG	TTCTTCCTCA	TCTCCTTTTG	GAAAAATTCT	26580
ACTCCATGCA	ATAGAAGTAC	GAAAAACATT	AAAGCCCATT	TCAGAAAACA	AGGATATATC	26640
TTCTTTATAT	TTATGATAAA	AATCAATACC	TATCAATTTT	AAGTTATCTT	CTGTAGGATT	26700

955

TTCTGTTGCT	TCTCCTAATC	CACCTTTGGG	TAACACATCC	TGAACTGATA	AGCCCTTACC	26760
ATCTTCATTA	TATGCTCCCT	CTACTTGATT	AGCTGCAACA	GCTCCACCCC	AAAGAAAATC	26820
ATCTGGAAAA	ATGGTCATAA	CTTTCCTCCA	TTATAATATT	ACCAGTAATF	CCTTAGAATG	26880
CTCGATTGTC	TGATTATTAG	GTAATACTAA	TACATCTAGA	AAATCATTGG	TATTCGTTAC	26940
AATTACTGGT	GTAAGTGTTC	CGTAGCCTTT	AGTCTTGATT	AAATTCAGT	CCATTTCAA	27000
AATCAACTGA	TTTTTGAAAA	CTCTGTCTCC	TTCTTCTACA	TGACTAATAA	AACCTTGACC	27060
TTTTAGCTCA	ACAGTATCTA	ATCCAATATG	AATTAGTAAC	TCAACACCCT	CATCACTCTT	27120
CAATCCAATT	GCGTGCTTAG	TCGGAAAAAT	ATTTGTAATT	TCCCATCAA	ATGGTGCATA	27180
AACCTTACCT	TCACTTGGGA	TAATCGCTAC	TCCGTCTCCA	ATTAGTTTAT	CTGAAAAATG	27240
TTTATCCTGG	ACATCGCTTA	ACGGAATGAT	TTCTCCTGAT	ATAGGAGAAA	ATATCATTTT	27300
TTTATTTGAA	ACTCCAGCTT	CAACTTCTAA	ATFGCTAGAA	CTCTCTTCTT	CATCGATTCC	27360
AAATATATAA	GCTAATACAA	AGGTAATAAC	AACCGAAATG	ACCGCCACAA	TTAAAGCATT	27420
TACAATATTT	GATGGCACAT	CAGAATAAAT	AAATTGAGGC	AACGCTATCA	AAGATGGGAC	27480
AGCAAATAGA	TATGCTTTAA	CACTAGTAAG	ACCTGCAAAT	AATCCCGCTA	ATCCACCACC	27540
AATCATAGCT	GCATAAAGCG	GTTTTTTATA	TTTTAAAGTC	ACACCATATA	ATGCAGGTTT	27600
GGTAATCCCT	GCAAGTAAGG	CTGAGAAACC	TGCTGCAAAA	GCAATTTGTT	TTGTATTATT	27660
ATTTTTACTC	TTTAATGCAA	CAGCCATCGA	AGCAGCCCCT	TGAGCTAAGT	TTGACCCTAA	27720
CATTGCTGGA	AGAATTAATA	CGTCTGGAGT	AGCAATAGAT	GCCGCCAAAA	AAATAGGTGC	27780
AAAAGCCCAA	TGCATTCCAG	TCATAACAAT	AAATGGCATA	ATAGCACCAA	GAATAGCTAA	27840
TGTAAGCCAT	CCAGCTACAC	CATACATTTG	CCCAACTAGA	TTTGATAATC	CTTACCACAC	27900
AATTACTCCA	ATAGGTCCGA	CTACAACATA	GGCAATACAG	CTTGATACTA	ATAATACTAG	27960
CGTAGGTTGC	AAAAAACTCT	TAGTAATAGC	TAGTGTAAAT	TTAGCAATTA	TTTTTTCAAT	28020
ATATTTTATC	AACCAAACCA	TAATAAGAAT	TGGAACGACT	GATGAACCAT	AACTAGCTGG	28080
TGTCACAGGT	GCACCAAATA	AACTAAGAGG	ATCCCTGAT	TGCACCATT	GAACAAAATT	28140
TGGATGGAGA	AGTACACCTG	CTACAGACAT	AGCTAATGTA	GATGTTACTT	TTAATTTTTG	28200
TGATGCAGAA	TAAGCTAATA	ACAGCGGTAA	GAAATAATAT	GGAGCATCCC	CAAAAAATGT	28260
CAAAAAGCA	ATAGTCTGAG	AATCTGATTG	CAATATACCA	AGCATTGGTA	AAATGATTAC	28320
CAAGACTTTC	AACATACCTC	CCCCTAACAT	TGCTGGAATG	ATTGGAGTCA	TGGAACCAGC	28380
GATATACTCA	ATGATTCTTT	CTAAAAATAT	CCCTTTGTGC	CCTTGAACAA	CTGAATCGGA	28440

956

TTCAAAATTG CCAAGTTTAA CGAATTCCTT ATAATAATTA GCTACATCAT TACCAAGTAT	28500
AATTTGATAT TGTCCATTCT TTTTCATAAT ACCTATTACA CCTGGTATCT TCTTCACATC	28560
ATCATCATTG ACTAAATTTT CATCTTTTAA TTCTAATCTT AAACGTGTTA CACAATGGGT	28620
AACTCTATTG ACATTTTTTT CACCTCCAAT TACATCGAGG ATTTTTTGTA CCGTATCTTT	28680
ATAAATCATG GTATTCTCCT ATTCTATTAA TCTAAATTTT TTGTTAAGCG ACGAATATGA	28740
GCCATCAAAT AAACTAATTC ACTAGAAGTC AGCAAATAAT TGTACTCCGT TTGTATAAAC	28800
ATTGCTACCT GTTCACCACA TTCATATTCT CTAGGATATT TATTTTTCAT TAATGCTAAC	28860
AAGTCTTCAT CATCATCGTC GG	28882

(2) INFORMATION FOR SEQ ID NO: 141:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 12835 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 141:

GCCTATGTCT TTTTCAAAAA AATGCTTGAC TTGAGACGGG AACTAGGGAA GTCTAAAGGC	60
GGAAGGCATT GATTTATACT CTTCGAAAAT CTCTTCAAAC CACGTCAACG TCGCCTTGGA	120
TTATATATGT AACTGACTTC GTCGATGCTT ATCTACAACC TCAAAGCAGT GCTTTGAGCA	180
ACTTGCGGCT AGTTTCCTAG TTTGCTCTTT GATTTTCATT GAGTATTATA TTACTTTCTA	240
TTTGTAGGAG GTGGCTTATG AAGATTCCTC TCTTAACTTT TGCAAGGCAT AAATTTGTTT	300
ATGICTTGCT TACTTTGCTT TTTCTTGCTT TGGTTTATCG TGATGTTTTG ATGACTTATT	360
TCTTTTTTGA TATTCATGCG CCCGATCTAG CTAATTCGA TGGACAAGCA ATTAAAAATG	420
ACTTATTAAA ATCAGCATTG GATTTTCGTA TTCTCCAGTT CAATCTAGGT TTTTATCAAT	480
CATTTATTAT TCCAATCATC ATTGTTTTGC TAGGTTTTCA ATATATTGAG CTGAAAAATA	540
AAGTTTTACG ATTGAGTATT GGAAGAGAAG TGAGTTATCA AGGGTTAAAA AGAAAGTTGA	600
CTTTGCAAGT TGCAAGTATC CCTTGTTTGA TATATTTAGT GACTGTGCTG ATAATTGCAA	660
TTATAACCTA TTTCTTTGGG ACTTTTTCTC CTCTTGATG GAATCTCTA TTTTCTGATG	720
GAAGTGGTTT ACAAAGACTC CTAGATGGAG AGATAAAAAG CTATTTGTTC TTTACTTGTG	780
TCCTACTAAT CGGTATTTTC ATCAATGCAA TCTATTTTTT ACAAATAGTT GATTATGTGG	840
GGAATGTGAC TCGTTCGGCA ATCACCTATT TGATGTTTCT TTGGCTTGGT TCTATGCTGC	900
TTTATAGTGC CTTCCTTAC TATATGGTTC CTATGACGAG TTTGATGCAA GCTAGCTATG	960

GGGATGTAAG TTTGATGAAA CTCTTTACTC CTTATATCCT TTATATTGTC CCTTACATGG	1020
TGCTTGAAAA ATATGAAGAT AATGTTTAAAG AATTTTAAACA ATATTTTGCT AAATAGAAAAG	1080
ATTGTTTTTAC TACTTCGTAT AGTTCGTGATG ATGATTTTGA TAAACCATCT ATTGTCAACA	1140
GCGGTTCAAAA AGCAGGATGC TGTATCTTT TTCAAGAGAG AATGATTTC AATTTTTTCC	1200
TATAATGACT ATTCTGAAGC GAATTTAGAA ATCCCAAAC TATGTTAAA CCTTTCGCTT	1260
TTCATGGTAG GATGGCTCTC TGTCATTTTA CTTGAAAGTG ATTTGGCAGA CCATTACCAT	1320
CACTTGATTC GCTATCAATC AAGCTCCTTT TTCGATTATA CAAGGAAACG ATTTGGTTGTC	1380
ATTTCTAAAT TTTTACTCA AGATTTGTTT GTCTGGTTTC TTGGTTTACT TCCTCTAGGA	1440
ATTCATTTCA AACAGTCGC ACTTTCTTT TTACTIONGCTC AGTTAATGAT GTTGTACTTA	1500
CTACTGTCTT ATCTGATAGC ACTGATTAGT GCGGGCGCTG GTTTTTCCTT TTTTCTCTAT	1560
TTTTTAGCAT TTGTGGGACA AGAATGGATG ATGGATCATA TTGTAACAGT GTATTTAGTA	1620
CTCTTAAGTT TATTAGTTAT GTTGATTGTT AGTCGCTTGG AAGAGAAAAT TAAGAAAGGA	1680
TAAACGATGA GACTTGAAAT TATAAATGGA CAGAAAATTT ATGGGAAAAG ACCTATTTTA	1740
AATCAGTTGA ATTTGGTGT TCAATCAGGA AAAATTTATG GACTTAAAGG TGATAATGGA	1800
TCTGGCAAGA CGGTTCTTTT AAAGATACTT GCTGGTTATA TTAAGCTTGA CAAAGGAAAA	1860
GTTCCTCAAG ATGGTAAAGT TTACGGGGTA AAAAATCATT ATATTCAGGA TGCAGGAATT	1920
TTAATGAAA AAGTCGAGTT TTTATCTCAT TTATCCCTGA GAGAAAATTT GGAACGTTA	1980
AGGTATTTTT CATCTAAAGT TACGGAAAAA AGAATTCCTT ATTTGGATTCA ATACTATGAT	2040
TTACAGGAAT TTGAAGACAT TGAATACCGT CATTTATCCT TAGGAACAAA GCAAAAAATG	2100
GCCTTGATTC AAGCCTTAT TTCTCTCCT TCTATACTCT TTCTCGATGA ACCTATGAAT	2160
GCTTTGGATG AGAAGAGTGT GAGGTTAACC AAACAGGTCA TTTTATCTTA CCTGAAAAAA	2220
GAAAATGGTC TGGTTATCCT GACGTCGCAC ATATCGGAAG ATATTTCAGA CCTTTGTACA	2280
GATGTATTAG TTGTCGAAAA TGGACATATA CAAATGTAAA GGATATACAA TCCTAGGAGA	2340
TGGCTTATGG CACATCTAAA ATCATTATTT ACACGATATT CCAAGGTTTA TATTGGTTTA	2400
GTTCCTGCTGA TCTGGCTGTC TTTCTTCTTT ATCCCTTGGG ATAAACCACT TCTGGGGATA	2460
AGGATTGACA TCTTCATCAT ACAGAAAAATC TTGCTAGCTT TTGGAATTC GTCCATTCTC	2520
ATGGCCTTGC TGTCCAAGAA AGTCAGTCTC TTTGTTTTTG GACTGATTG CTGTCTTTCT	2580
CFTTGGATTA ACTTATTTAT CACATTTGCC ATTTTGCCGA TTTTGGCAA TTAAACAGTC	2640
ATAAAAGTCG GAGAGGTTAG CTTGAAAATC AACCTCTTTT TCCTTTTCAA AATGGGGATT	2700

CTTCCTTGAA	AATAATCAGT	AATTGTGCTA	AAATTAAAGG	AACATTCTAA	AATATTCGGA	2760
ATTTAAAGTA	AGGAAAAACA	TGGCTAATAT	TTTAAAAACA	ATTATCGAAA	ATGATAAAGG	2820
AGAAATCCGT	CGTCTGGAAA	AGATGGCTGA	CAAGGTTTTT	AAATACGAAG	ACCAAATGGC	2880
TGCTTTGACT	GACGACCAAC	TAAAAGCAAA	AACAGTTGAA	TTTAAGGAAC	GTTATCAAAA	2940
TGGAGAATCA	CTGGATTCAT	TGCTTTACGA	AGCATTTCGG	GTTGTCCGTG	AAGGTGCCAA	3000
ACGTGTCCTA	GGTCTCTTCC	CTTATAAGGT	TCAGGTCATG	GGGGGGATTG	TTCTTCACCA	3060
TGGTGACGTG	CCAGAGATGC	GTACAGGGGA	AGGAAAACC	TTGACTGCGA	CCATGCCGGT	3120
ATACCTCAAT	GCCCTTTCAG	GTAAAGGGGT	TCACGTAGTT	ACGGTTAATG	AATACCTGTC	3180
AGAACGTGAC	GCGACTGAGA	TGGGTGAATT	GTACTCTTGG	CTTGGTTTGT	CAGTAGGGAT	3240
TAACCTGGCT	ACCAAATCTC	CAATGGAGAA	AAAAGAAGCC	TATGAGTGTG	ATATTACTTA	3300
CTCAACTAAC	TCAGAAATCG	GATTTGACTA	CCTTCGTGAC	AACATGGTCG	TTCGCGCCGA	3360
AAACATGGTA	CAACGTCCGC	TTAACTATGC	CTTGGTCGAT	GAGGTTGACT	CTATCTTGAT	3420
TGACGAGGCT	CGTACACCTT	TGATTGTATC	AGGTGCCAAT	GCGGTTGAAA	CCAGTCAGTT	3480
GTATCACATG	GCAGACCCTT	ATGTAAAATC	TTTGAACAAA	GATGACTACA	TCATCGATGT	3540
GCACTCTAAG	ACTATTGGTT	TGCTCGATT	AGGGATTGAC	AGGGCTGAAA	GCTACTTCAA	3600
ACTTGAAAAC	CTCTATGACA	TCGAAAACGT	GGCTTTGACT	CACTTTATCG	ATAACGCCCT	3660
TCGTGCCAAC	TACATCATGC	TTCTCGATAT	TGACTATGTG	GTGAGCGAAG	AGCAAGAAAT	3720
CTTGATTGTC	GACCAATTTA	CAGGTCGTAC	CATGGAAGGT	CGTCGTTATT	CTGATGGATT	3780
GCACCAAGCT	ATTGAAGCCA	AAGAAGGTGT	GCCAATCCAG	GATGAAACCA	AGACATCTGC	3840
CTCAATCACG	TACCAAAACC	TCTTCCGTAT	GTACAAGAAA	TTGTCTGGTA	TGACGGGTAC	3900
AGGTAAGACT	GAGGAAGAAG	AATCCCGTGA	AATCTACAAC	ATTCGTGTTA	TTCCAATCCC	3960
AACAAACCGT	CCTGTTCAAC	GTATTGACCA	CTCAGACCTT	CTTTATGCAA	GTATCGAATC	4020
TAAGTTTAAA	GCGGTTGTCG	AAGACGTTAA	GGCTCGTTAC	CAAAAGGGTC	AACCTGTCTT	4080
GGTTGGTACA	GTAGCGGTTG	AACTAGTGA	CTACATTTCT	AAGAAATTGG	TTGCAGCTGG	4140
TGTTCCCTCAC	GAAGTCTTGA	ATGCCAAAAA	CCACTATAGA	GAAGCCCAA	TCATCATGAA	4200
TGCTGGTCAA	CGTGGTGCCG	TTACCATCGC	AACCAACATG	GCGGGTCGTG	GTACCGACAT	4260
CAAGCTTGGT	GAAGGTGTTC	GTGAACTTGG	AGGACTTTGT	GTTATTGGTA	CAGAACGTCA	4320
TGAAAGTCCG	CGTATCGATA	ACCAGCTTCG	TGGACGTTCA	GGTCGTCAAG	GAGATCCAGG	4380
TGAGTCACAA	TTCTACCTAT	CTCTTGAAGA	TGATTTGATG	AAACGTTTTG	GTTCTGAACG	4440
CTTGAAGGGA	ATCTTTGAAC	GCTTGAACAT	GTCTGAAGAG	GCCATTGAGT	CTCGCATGTT	4500

GACGCGTCAG GTTGAAGCAG CTCAGAAACG TGTCGAAGGA AATAACTACG ATACCCGTAA 4560
 ACAAGTCCTT CAATACGATG ATGTCATGCG TGAACAACGT GAGATTATCT ATGCTCAACG 4620
 TTACGATGTC ATCACTGCAG ATCGTGACTT GGCACCTGAA ATTCAGTCTA TGATCAAACG 4680
 CACGATTGAA CGTGTCGTG ATGGTCATGC GCGTGCCAAA CAAGATGAAA AACTAGAGGC 4740
 AATTTTGAAC TTTGCTAAGT ACAACTTGCT TCCTGAAGAT TCTATTACGA TGGAAAGACTT 4800
 GTCAGGCTTG TCTGATAAGG CCATCAAGGA AGAGCTTTTC CAACGTTCCCT TGAAGGTTTA 4860
 CGATAGTCAG GTTTCAAAAC TACGCGATGA AGAAGCAGTT AAAGAATTCC AAAAAGTTTT 4920
 GATTCTACGA GTGGTGGATA ACAAGTGGAC AGATCATATC GATGCCCTTG ATCAATTGCC 4980
 TAACGCGGTT GGACTIONCTG GCTATGCTCA GAACAACCCT GTTGTGAGT ATCAGGCAGA 5040
 AGGTTTCCGT ATGTTTAATG ATATGATTGG TTCGATTGAG TTTGATGTGA CACGCTTGAT 5100
 GATGAAAGCA CAAATTCATG AACAAAGAAAG ACCACAGGCA GAACGTCATA TCAGTACAAC 5160
 AGCGACTCGC AATATCGCTG CTCACCAAGC AAGTATGCCA GAAGATTTGG ATTTGAGCCA 5220
 GATTGGACGC AATGAAC TTT GCCCATGTGG TTTCTGGTAAG AAATTTAAAA ACTGTCACGG 5280
 TAAAAGACAA TAAAATGAGA TAGTTTAGAG GCGGATATCT TGTGAAAAGT AAATTTTAC 5340
 TGGGTATCCG TTTGCTTTAT AAGGAGATGA GTTATGGTAT TTACAGCAA AAGCTCTAAA 5400
 ATAAATATAG AAGAAGTTCG TGCTTGTCA AAATTAGAAG GTCAGGCTTT GGAGAGGAAA 5460
 TCACAGCGAG ATCAAGAGCT AGAAGCCATT ATACGTGGAG AAGACCAGCG AATCTCTTG 5520
 GTAATCGGGC CATGCTCATC TGACAACGAA GAAGCTGTCC TTGAATACGC TAAGCGTTTG 5580
 GCAGTCCTAC AAGAAGAAGT GGCAGATCGT ATCTTTATGG TTATGCGTGT TTATACTGCC 5640
 AAACCCCGTA CCAACGGAGA TGCTATAAG GGCTTGATTC ACCAGCCTAA CGCGACAGAA 5700
 GCGCCTAGTC TTATCAATGG AATCAAAGCC GTTCGCCATC TTCACTATCG TGTCATCACA 5760
 GAAACAGGGA TGACAAC TGC TGATGAAATG CTTTATCCTG AAAACCTTCC GCTTGACAT 5820
 GATTTGATTT CTTACATGGC AGTTGGTGCC CGTTCAGTTG AAGACCAGCA ACACCGCTTT 5880
 GTGGCAAGTG GGGCAGGATT TTCTACTGGT TTTAAAAATC CAACCTCTGG AAATCTCAAT 5940
 GTCATGTTTA ATGGGATTTA TGCTGCTCAA AACAAACAAA GTTTCCTTTT CTTAGGAAAA 6000
 GAAGTAGAAA CAACTGGGAA CCCGCTTCA CACGCTATTC TTCGTGGTGC TCTTAATGAG 6060
 TATGGAAAAA ATATTCCCAA CTACTATTAT GACAATTTAA TTGATACCAT TGCCAGTAT 6120
 GAGAAAATGG GCTTGAAAA TCCTTTTATC ATCATTGATA CCAATCATGA CAATTCTGGT 6180
 AAGCAGTATA TTGAACAGAT CCGAATTGTC CGCCAGACCT TGATTAACCG TGCTTGAAT 6240

960

GAAAAAATTA AGCAGTTCGT TCGTGGTTTT ATGATTGAGT CTTATCTGGA AGATGGTCGA 6300
 CAAAATGAGC CAGAAGTATT TGGTAAGTCT ATCACAGACC CTTGCCTGGG TTGGGATAAC 6360
 ACAGAAGCTC TTGTCAGAGA AATTTACAAA ACGTTAGGAG AATAAGATGG CATTATTGGA 6420
 AAAAGGTCAA GAAATCGATA TGGAAATCAT CAAGGCTGAA ACCCAATTGT CTGCGGAAGC 6480
 CTTGAGACTC AAGGAAAGCC GTGACAGGGA ATTGGCAGAT ATTATTTTCAG GGAAGATGA 6540
 CCGTATTCTC TTGGTGATTG GTCCTTGCTC TTCTGATAAT GAAGAGGCGG TCTTGAATA 6600
 TGCTCGCCGT TTATCTGCCT TGCAAAAGAA GGTAGCGGAT AAGATTTTCA TGGTCATGCG 6660
 CGTGTATACT GCTAAGCCTC GTACCAATGG AGACGGCTAT AAAGGATTAG TTCACCAGCC 6720
 AGATACTTCT AAGGCTCCAA GCCTGATTAA TGGCTTGCAG GCTGTGCGCC AGTTGCACTA 6780
 CCGCGTGATT ACAGAGACTG GTTTGACAAC GGCAGATGAG ATGCTTTATC CGTCAAATCT 6840
 GATCTTGGTG GATGACTTGG TCAGTACCA TGCCGTTGGA GCTCGTTCTG TGAAGACCA 6900
 AGAGCACCGC TTTGTGGCTT CTGGGATTGA TGCACCAGTA GGGATGAAAA ATCCAACCTC 6960
 AGGAAATTTG GGTGTATATG TTAACGCCAT CTATGCTGCT CAAAACAAGC AAACCTTCCT 7020
 TTATCATGGG CAGGAAGTTG AGACATCAGG TAATCCTTTG GCCCATGTTA TCCTCCGTGG 7080
 AGCAGTCAAC GAGTATGGCA ATTATATGCC GAATTACTAC TATGAAAATC TACTCCAAGC 7140
 CATTGAACGC TATGAAACCA TGGGACTTGA AAATCCTTTT ATCCTCATTTG ACACCAACCA 7200
 TGATAACTCA GGCAAGCAAT ATATGGAGCA GATTCGAATT GTTCGCCAGA CCTTGCAGAA 7260
 TCGTGATTGG AATGAGAAAA TTAAAAAGAC GGTTCGAGGA TTTATGATTG AATCTTACCT 7320
 AGCAGATGGT CGTCAAAACC AACCAGAGAT CTTTGGTTGC TCTATFACTG ACCCTTGCCT 7380
 AGGTGGGAA AATACAGAGG CCTTGGTAGA AGAGATTTAT GTTACCTTGA CAAAATAAGT 7440
 GAAAAGGATG GAGTTGGGGA ATCTCAACTC CTTTTGATGA GAATGATAGT TGGACACGGA 7500
 ATTGACATCG AAGAATTGGC TTCGATAGAA AGCGCAGTTA CACGACATGA AGGATTTGCT 7560
 AAGCGTGAC TGACCCTCA GGAATGGAG CGCTTCACCA GTCTCAAAGG ACGCAGGCAA 7620
 ATAGAATATT TAGCTGGTCG CTGGTCGGCT AAGGAGGCTT TTTCCAAGGC TATGGGAACG 7680
 GGCATTAGCA AGCTCGGTTT TCAGGATTTG GAAGTCTTGA ACAATGAACG TGGGGCGCCT 7740
 TATTTTAGTC AGGCACCATT TTCAGGAAAG ATTTGGCTGT CTATCAGCCA CACCGATCAG 7800
 TTTGTGACAG CCAGTGTCAT TTTGGAGGAA AATCATGAAA GCTAGTCCAC ATAGACCAAC 7860
 CAAGGCTCTG ATTCATCTGG GAGCTATTCTG ACAAATATT CAGCAAATGG GGGCTCATAT 7920
 CCCTCAAGGA ACGCTCAAGT TGGCTGTGGT TAAGGCCAAT GCTTATGGTC ATGGAGCTGT 7980
 TGCCGTTGCC AAGGCAATTC AAGATGATGT TGATGGCTTT TGCGTTTCCA ATATCGATGA 8040

AGCCATTGAA CTCAGACAAG CTGGACTCAG CAAGCCAATC CTCATTTTAG GAGTTTCTGA 8100
AATCGAAGCT GTTGCTCTAG CTAAGAATA TGACTTCACC TTGACAGTGG CTGGACTGGA 8160
GTGGATTCAA GCACTCTTAG ATAAGGAAGT GGACCTAACT GGATTGACAG TCCACCTCAA 8220
GATTGATTCA GGGATGGGAC GGATTGGTTF TAGAGAGGCA AGTGAGGTTG AGCAGGCTCA 8280
AGATTTGCTC CAACAACACG GTGTTTGTGT TGAAGGAATC TTTACCCACT TTGCTACTGC 8340
TGATGAGGAA TCAGATGACT ATTTAATGC CCAGTTAGAA CGGTTTAAAA CTATTTTAGC 8400
TAGTATGAAG GAAGTCCAG AGCTGGTTCA TGCTAGCAAT TCTGCAACGA CTCTTTGGCA 8460
TGTAGAGACT ATTTCAATG CCGTTCGTAT GGGAGATGCC ATGTATGGCC TCAATCCAAG 8520
TGGAGCGGTC TTGGATTTGC CTTATGATTF GATACCGGCC TTGACCTTGG AGTCTGCTCT 8580
GGTTCATGTC AAGACAGTTC CAGCTGGAGC TTGCATGGGC TATGGAGCAA CTTATCAAGC 8640
GGATAGCGAG CAAGTCATCG CGACCGTCC AATCGGGTAT GCAGATGGAT GGACAAGAGA 8700
CATGCAAAAT TTCTCTGTCT TGGTAGATGG CCAAGCTTGC CCAATTGTCG GCAGGGTTTC 8760
GATGGACCAA ATCACTATTC GATTGCCTAA GCTTTATCCG CTAGGAACCA AGGTAACCTT 8820
GATTGGCTCC AATGGGGATA AGGAAATCAC TGCAACTCAG GTAGCGACCT ACCGCGTAAC 8880
CATTAACTAT GAGGTGGTTF GCCTCCTCAG CGACCGTATT CCGAGAGAAT ATTATTAGAA 8940
AAGAAAGGAG TGGAGCATGA ATCTACATCA ACCCTTGCAT GTCTTGCCTG GTGTGGGACC 9000
AAAGTCAGCA GAAAAATACG CCAAACCTAGG AATTGAAAAC TTGCAAGATC TCTTGCTCTA 9060
CTTTCCTTTC CGTTATGAAG ACTTCAAAAC CAAGCAGGTG CTGGAGCTGG AAGACGGTGA 9120
GAAGGCAGTT CTTTCTGGTC AGGTAGTGAC TCCTGCTAGT GTCCAGTATT ATGGTTTCAA 9180
GCGCAATCGC CTGCGTTTFA GTCTCAAGCA GGGAGAGGTC GTTTTGGCGG TGAATTTCTT 9240
TAACCAGCCC TATCTGGCTG ATAAAATAGA GTTGGGAGCA ACCCTTGCTG TCTTTGGAAA 9300
ATGGGACCGC GCTAAGGCTA GTCTGACTGG GATGAAGGTT CTGGCTCAGG TAGAAGATGA 9360
CCTCCAGCCT GTCTATCGTC TGGCTCAGG AATCAGTCAG GCCAGTCTGG TCAAGGTCAT 9420
CAAGACGGCT TTGATCAGG GACTGGACCT CTTGATAGAA GAAAATCTGC CCCAGTCTTT 9480
ACTAGACAAA TACAACTCA TGTCCTGTTG TCAGGCAGTC CGTGCTATGC ATTTTCCAAA 9540
GTATTTGGCA GAATACAAGC AGGCTCTTCG CCGTATAAAG TTTGAGGAAC TCTTTTATTT 9600
CCAAATGCAG CTGCAGATGC TCAAGTCTGA AAATAGAGTT CAGGGAAGTG GTCTGGTTCT 9660
GAATTGCTCT CAGGAAAAAG TGACAGCAGT TAAAGTAAGT CTTCTTTTGG CCCTGACCCA 9720
AGCTCAGGAA AAGAGTTTGC AGGAAATTTT AACTGATATG AAGTCCGACC ACCACATGAA 9780

962

TCGTCTCCTA	CAAGGGGATG	TGGGGAGTGG	AAAAACGGTA	GTCGCTGGCT	TGGCCATGTT	9840
TGCGGCAGTG	ACAGCAGGTT	ATCAGGCTGC	CCTAATGGTA	CCAACAGAAA	TCCTCGCAGA	9900
GCAACACTTT	GAGAGTTTAC	AGAACCTTTT	TCCCAATTG	AAACTGGCTC	TCTTGACAGG	9960
TTCTTGAAA	GCTGCAGAAA	AGAGAGAAGT	CTTGGAGACC	ATTGCCAAGG	GTGAGGCTGA	10020
TTTGATTATA	GGAACCTCAC	CTCTGATACA	AGATGGGGTG	GAGTATGCTC	GTCTTGGTTT	10080
GATTATTATC	GATGAGCAGC	ACCGTTTTGG	TGTAGGGCAA	AGGCGTATTT	TACGGGAAAA	10140
AGGTGACAAAT	CCAGATGTCC	TCATGATGAC	GGCGACTCCC	ATCCACGGA	CGCTTGCCAT	10200
CACAGCCTTT	GGAGATATGG	ATGTTTCCAT	TATCGACCAG	ATGCCAGCAG	GTCCGAAGCC	10260
TATTGTGACG	CGCTGGATCA	AACATGAGCA	ACTACCTCAG	GTCTTGACTT	GGTTAGAGGG	10320
GGAAATTCAA	AAAGGTTCCT	AAGTCTATGT	CATCTCTCCT	TTGATTGAAG	AATCAGAAGC	10380
TCTAGATTTG	AAAAATGCCA	TTGCCTTATC	AGAGGAGTTG	ACGACTCATT	TTGCAGGCAA	10440
GGCAGAGGTG	GCTCTTCTAC	ATGGTAGGAT	GAAGAGTGAC	GAAAAAGACC	AGATCATGCA	10500
GGATTTCAAG	GAGAGAAAGA	CGGATATTCT	GGTTTCGACG	ACGGTTATTG	AGGTTGGGGT	10560
CAACGTTCCT	AATGCGACTG	TCATGATTAT	CATGGATGCC	GATCGCTTCG	GTCTCAGTCA	10620
ACTTCACCAG	CTTAGAGGTC	GTGTCGGTCG	GGGGGACAAG	CAGTCCTACG	CTGTTCTCGT	10680
TGCTAATCCC	AAGACGGATT	CTGGGAAAGA	CCGCATGCGT	ATCATGACAG	AAACGACCAA	10740
TGGATTTGTC	CTTGCGGAGG	AAGATTTGAA	AATGCGTGGT	TCTGGTGAGA	TTTTTGGAAC	10800
CAGACAGTCA	GGACTTCCAG	AGTTCCAAGT	GGCTGATATT	ATCGAAGATT	TTCCGATTTT	10860
AGAAGAAGCA	AGAAAGGTTG	CTAGCTACAT	TAGTTCTATA	GAAGCTTGGC	AAGAAGATCC	10920
AGAGTGGCGC	ATGATTGCCC	TTCATCTGGA	AAAGAAAGAA	CATCTGGATT	AAGCTTTCTC	10980
TAAGGAAAAC	TTATACTCAA	TGAAAATCAA	AGAGCAAACCT	AGGAAGCTAA	CCGCAGGTTG	11040
CTCAAAACAC	TGTTTTGAGG	TTGTGGATGA	AACTGACGAA	GTCAGCTCAA	AACACCGTTT	11100
TGAGGTGGCA	GATAGAACTG	ACGAAGTCAG	TAACATATAT	ATACGGTAAG	GCGACGCTGA	11160
CGTGGTTTGA	AGAGATTTTC	GAAGAGTATT	AAGCTAGTTT	TTAGGTTTGG	CTCTTATACT	11220
AGAGTCATCA	AAAAGAAACG	AGGACTCTCA	TATGACAGTA	ACGATTAAAG	TAAATTACCA	11280
AACCACTTTC	CAAAAGAAGG	AAGCAAAAAA	CTAGTATAAA	CAGAAGAGAG	AGCGAAATGC	11340
TCTTTTTCG	TTTCTAAAAC	TACTTTCAGC	CCATCATCCT	AAAAGTAAAG	AATCTAAATT	11400
CACTTTCAT	TTACCCTTCT	TTCTTGCAAT	GATTACATAG	ATATGCTACA	GTTGTGGTAA	11460
CGATTACAAA	ATAAAAGGAG	CATGCTATGA	AAAATCCAGC	TTTGCTAGAA	GAAATTAAGA	11520
CCTATAGAGG	AAGGGATGAG	GTTCCGGAAG	ACTTTGATGA	TTTCTGGGAT	GGGGAAGTGA	11580

AAAATGTTTC CACGCTTCCA TCCTACCACT TGGAGGAAAG AGATTTCCAC ATTCCTCAAG 11640
 TCAAGTGCTA TGAGTTAACA TTTGAAGGAA GCAAGGAAGG AAAGGTCTAT GCACGCATTG 11700
 TTCTTCCAAA GAGTGAGGAG AAGGTCCCAT TAATCTTCCA TTTTCATGGT TATATGGGAC 11760
 GTGGCTGGGA CTGGGCCGAC ATGCTGGGCT TCACCGTAGC TGGTTACGGT GTTGTTFCCA 11820
 TGGATGTGCG GGGCCAGTCA GGTTACTCAC AAGACGGCTT GCGTTCTCCT TTAGGAAATA 11880
 CCGTGAAGGG GCATATTATC CGTGGTGCTG TGAAGGTCG GGACCACCTC TTTTATAAGG 11940
 ATGTTTATCT GGATATTTAC CAGTTGGTCG AAATTGTTGC TAGTCTGTCT CAGGTTGATG 12000
 AGAAGCGTCT TTCTAGCTAT GGTGCCTCAC AAGGAGGGGC TCTAGCTCTA GTTGCAGCAG 12060
 CGCTCAATCC TCGAATTCAG AAAACAGTTG CCATTTATCC CTTCTTGTC AACTTCAGAC 12120
 GGGTGATTGA GATTGGTAAT ACTAGCGAGG CTTACGACGA ACTTTTCCGT TATTTCAAGT 12180
 TTCACGACCC CTTCCATGAA ACAGAGGAGG AAATCATGGC GACCCTTGCC TATATCGATG 12240
 TCAAAAATCT TGCCCATCGT ATCCAAGGTG AGGTTAAGAT GATTACGGGC TTGGACGACG 12300
 ATGTTTGCTA TCCCATTACC CAGTTTGCGA TTTATAATCG TCTGACCTGC GATAAAACCT 12360
 ATCGCATCAT GCCTGAGTAT GCTCACGAAG CCATGAATGT ATTTGTCAAT GACCAAGTCT 12420
 ACAACTGGCT CTGTGGAAGT GAGATTCCTT TTAAATATCT AAAATAAGGA GTCGACTCTA 12480
 AGCACAAAAT CTTAAAAATT ACAAACACGC ATAGTATCAG GGGATTAAGA AAACTTTATA 12540
 CTATGCGTTT TATCATGGAA ATATAGTAAA ATGAAATAAG AACAGGACAA ATCGATCAGG 12600
 ACAGTCAAAT CGATTTCTAA CAATGTTTTA GAAACAAATG TGTACTATTC TAGTCTCAAT 12660
 CTATTATATT TATAGAATTT TTTGTTGCTA GATTTGTCAA ATTGCTTAAA ATAATTTTTT 12720
 TCAGAAAGCA AAAGCCGATA CCTATCGAGT AGGGTAGTTC TTGCTATCGT CAGGCTTGTC 12780
 TGTAGGTGTT AATACTTTTC AAAAATCTCT TCAAACCACG TCAGCTTCGC CTTGC 12835

(2) INFORMATION FOR SEQ ID NO: 142:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 5020 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 142:

GGGGATATGA AGAACAAAAG AATATTTAAA GACTTCCAAG CTTCAAAAAT GAGTTTAAAC 60
 ATTTACACAA GCCCCTTGTT AGCCTTTGTT TTTGTCTTCA TAGGAGAGTT TGTGGCTTTT 120

964

ACTTTGTATG GTATTGGCTT GTTAGCTCTC ATCGGACTTG CTAGAAATTT TGGAGAGGCT	180
GGTCAAAATC TTGCAAGCTA CTTGCAGACC TTGCATCAGA GCTTGACGGA TAAAACAAGT	240
GACTTTCGTT TAATTTTAGG ATTACTGGCC TTTGGTTATT CTTAACACTG TGTTCAGATG	300
GACAAGAAAA GTTGAGAAAA GACCTATTCG AACCTTGGGA TTTTATAGAG AGAATTTCTT	360
CAGCAATCTT CTGAAAGGAT TTAGTCTAGG CCTGGCACTT TTTCTTCTGA CCTTGTTAGG	420
TTTAGTGGTC TTAGGTCAAT ATCGTTTGGA ATCCATTCAC TTGAATCCTT ATTCTCTTGC	480
CTTTGTCGTC TTTACTATCC CATTTTGGAT TTTACAGGGG ACAGCAGAAG AAGTGGTGGC	540
CCGTGCTTGG CTTACTTCCTC AATTGGCCTC AAGAACCAAT CTAAAAC TAG CTATTCTTAT	600
ATCTAGCCTG TTTCTTTACCC TGCTTCATAT GGGCAATTCT GGCTCACCCT CTCTATCTCT	660
AGTAAATCTC TTTTATTCG GAGTTGCCAT GGCTCTTAC CTTCTCAAAA CTGATACAGT	720
TTGGGGTGTT GCAGGTATTC ATGGTGCTTG GAATTTTGCT CAGGGTAATC TCTTTGGGAT	780
TTTAGTTAGT GGTCAACCGT CAGaACGTCT CTGATGACCT TTTTACCACA AGGCAATCAA	840
GATTGGCTAT CAGGTGGTTC TTTTGGCATA GAAGGTCCA TTATGACAAG TCTGGTATTA	900
CTACTGCTGA TTGTCTATCT TGCTAATAAA TTAAAGAAAG AAAATGAAAG GATGTGACTT	960
CGGTCCGTCC TTTTCTTCGT GAAAATACTA TAAGTATGCT AAAATAGGAA TAGCACATGG	1020
AGAGAGGATT CTTATGATCA ATCACATTAC AGATAATCAA TTTAAACTAG TATCAAAATA	1080
TCAACCATCA GGAGATCAAC CCAAGCTAT CGAGCAGTTG GTGGATAACA TTGAGGGGGG	1140
AGAAAAAGCT CAGATTCTGA TGGGGGCGAC TGAACAGGG AAGACCTATA CTATGAGTCA	1200
GGTCATTTCT AAAGTCAATA AACCAACTCT GGTTATTGCC CACAATAAAA CTCTGGCTGG	1260
TCAGCTCTAT GGGGAGTTA AGGAATTTTT CCCTGAAAAT GCAGTTGAGT ATTTTCGTATC	1320
CTACTATGAT TATTACCAGC CAGAGGCCTA TGTCCCTTCT AGCGATACCT ATATTGAGAA	1380
GGATAGTTCT GTCAATGACG AGATTGACAA GCTTCGCCAC TCAGCTACCT CAGCCCTTTT	1440
GGAGCGTAAT GATGTTATTG TCGTGGCCTC AGTCTCTTGT ATCTATGGTT TGGGTTCCGC	1500
CAAGGAATAC GCTGATAGTG TCGTTAGTCT CCGTCCTGGT CTAGAGATTT CTCGTGATAA	1560
ACTCTGAAT GACTTGGTCG ATATTGAGTT TGAACGTAAT GATATTGATT TCCAACGCGG	1620
AAGATTTTCG GTTCGTGGGG ATGTGGTAGA GATTTTCCCA GCTTCCCGAG ATGAACATGC	1680
CTTTCGAGTA GAATTTTTTG GAGACGAAAT TGACCGTATT CGTGAAGTTG AGGCTCTGAC	1740
AGGTCAGGTG TTGGGAGAAG TGGATCATTT AGCGATTTTC CCAGCGACAC ACTTTGTGAC	1800
CAATGACGAC CACATGGAAG TTGCCATTGC AAAGATTCAG GCCGAGTTGG AAGAACAATT	1860
AGCTGTCTTT GAAAAGGAAG GTAAACTGCT TGAAGCCCAG CGTTTGAAC AGCGGACAGA	1920

965

GTATGATATC	GAAATGTTGC	GTGAGATGGG	CTATACCAAT	GGGGTTGAAA	ATTAT'TCTCG	1980
CCACATGGAT	GGACGGAGCG	AAGGAGAGCC	TCCTTATACG	CTTCTCGACT	TCTTCCCAGA	2040
TGATTTCTTG	ATTATGATTG	ACGAGAGTCA	TATGACCATA	GGGCAAATCA	AGGGCATGTA	2100
CAATGGAGAC	CGTTCGCGTA	AAGAAATGCT	GGTTAATTAT	GGTTTCCGTT	TGCCGTCTGC	2160
TTTGGACAAT	CGTCCTCTCC	GTCGGGAGGA	GTTTGAGAGT	CACGTTCATC	AGATTGTTTA	2220
CGTTTCAGCG	ACACCTGGTG	ACTATGAAAA	TGAACAGACC	GAGACAGTGA	TTGAGCAAAT	2280
CATTCGTCCA	ACGGGACTCT	TGGATCCAGA	GGTGAAGTC	CGTCCGACTA	TGGGACAGAT	2340
TGATGACCTC	TTGGGTGAAA	TCAATGCCCG	CGTTGAAAAA	AATGAGCGTA	CCTTTATCAC	2400
AACTTTGACC	AAGAAAATGG	CAGAGGATTT	GACCGACTAC	TTCAAGGAAA	TGGGTATCAA	2460
GGTCAAGTAC	ATGCACTCGG	ATATCAAGAC	CTTGGAACGG	ACGGAGATTA	TCCGTGACCT	2520
GCGCTTGGGT	GTC'TT'GATG	TCTTGGTCGG	AATTAACCTG	CTCCGTGAAG	GAATTGACGT	2580
TCCTGAAGTG	AGCCTCGTAG	CTATTCTCGA	TGCTGACAAG	GAAGGTTTCC	TTCGCAACGA	2640
ACGTGGACTC	ATCCAGACCA	TTGGACGTGC	TGCACGTAAT	AGCGAAGGTC	ATGTTATCAT	2700
GTATGCGGAC	ACGGTTACCC	AGTCTATGCA	ACGTGCTATC	GATGAAACTG	CCCGCCGTCC	2760
CAAAATCCAG	ATGGCCTATA	ATGAAGAACA	TGGTATCGTT	CCACAAACCA	TCAAGAAAGA	2820
AATCCGTGAC	TTGATGCTG	TGACCAAGGC	AGTTGCTAAG	GAAGAAGACA	AGGAAGTCGA	2880
TATCAATAGC	CTCAACAAAC	AAGAGCGCAA	AGAACTAGTC	AAAAAGCTTG	AGAAACAAAT	2940
GCAAGAAGCA	GTTGAAGTGC	TTGACTTTGA	ACTAGCAGCT	CAGATTCGTG	ATATGATGCT	3000
GGAAGTCAAG	GCCTTGGATT	AGGGGAATAG	TATGATTTAT	TTAAGAAAGT	TAAAGAAAGA	3060
AGATTTGATG	TCTTTATGGG	AAATGGCTTA	TTCACAGCTT	AATCCAGTTT	GGAAACAGTA	3120
TGATGCTCCC	TATTATGATG	ATTATCAGTA	TTTTTCAAAT	TTTAAAGAAT	TCGAACTACA	3180
AAAATCAGAA	TCCATTTTAA	GCAACTCAA	TCGCCTGGT	ATTTTGTGTT	ATGATAAACT	3240
AGTTGGGACT	GTTTCGCGTT	ATTGGGTATG	TAAAGAAACA	AGATGGATGG	AATTGGGAAT	3300
TGGTATTTAT	GATAAAAAAT	TCTGGAACAC	TGGTATTGGG	AAAGTTGCTA	TGTTGCAGTG	3360
GATAGATAGG	ACGTTTCAGG	ATTACTTGGA	GTTGGAGCAT	CTGGGTTTGA	CAACTTGGTC	3420
AGGAAATATT	GGTATGATGA	AACTTGCTGA	AAAATTAAGA	ATGAAAAAAG	AAGCTCATAT	3480
TCCAAAAGTT	CGTTATTATC	AAGGTAAATA	TTTTGACAGT	ATTAAATATG	GTATTTTGAG	3540
AGAAGACTGG	GAGAAAATAA	ATGACGGTTA	TTATCAAATC	AATGGAAACT	CCTGAAGAGA	3600
TAGAAGGTAA	ATCCTTCGTT	CACTGGCAAA	CGTGGAGAGA	GGCTTATGAT	GACCTTTTGC	3660

966

CTGCGGAATT TCAGGAGACA ATGACATTAG AAAGATGTCG ACTCTTTAGT CAAAAGTATC 3720
 CAGAAAATAC ATTGATTGCG ATGGATGGTG TGAAGATAGT TGGTTTTATA AGTTATGGCA 3780
 ACTGTCTGTA TGAGACTATT CAAGCTGGTG AAATTATTGC TTTATATGTT TTAAGAACT 3840
 ATTATGGAAA AGGAATCGCA CAAAAGTTAG TGAAGCAGC TTTGACTGAT CTTAATCATT 3900
 TTTCTGAAAT TTTCTTATGG GTATTGAAAG ATAACAAGCG CGCCATTGCT TTCTATCAA 3960
 AAATGGGTTT TACTTTTGAT GGACAAGAAA AAATACTTGA ACTTGGAAG CCTATAAAGG 4020
 AAAAACGGAT GGTATTCTAT TCTAAATAAT TCTCAAAAGT AAAAGCTAAT ATGGTACCAA 4080
 GTCTGAAAAT TTAATAAATT AGAAAGCGAG TAAATTTATG TCCCCTTCCC AATTAACAAT 4140
 TTTAACAAT ATCTGTCTGA TTGAAGACCT CGAACTCAG CGCGTGGTGA TGCAGTATCG 4200
 CGCCCCTGAA AACAAATCGCT GGTCTGGTTA TGCCTTTCCT GGAGGTCATG TAGAAAATGA 4260
 TGAGGCTTTT GCGGAGTCTG TCATTCTGTA AATCTACGAA GAAACAGGGT TGACTATCCA 4320
 AAATCCTCAA CTTGTCTGCA TTAATAAATT GCCACTAGAT ACAGGTGGGC GCTATATTGT 4380
 CATTTGTTAT AAGGCGACTG AGTTCTCTGG TACCCTTCAA TCTTCAGAAG AGGGAGAAGT 4440
 TTCTTGGGTG CAAAAGACC AGATTCCAAA CTTAAATCTG GCCTATGATA TGCTACCATT 4500
 GATGGAAATG ATGGAAGCTC CCGACAAGTC AGAGTTTTTC TACCCTCGCC GTACAGAAGA 4560
 CGATTGGGAA AAGAAAATCT TCTAGTCTTT TACTAAATAA CCTAGCTGAT CCAAGGCCTC 4620
 CTCGATATAG TGGAGTCTT GTTGTGTCTC GGCTTCAACT AGGTGATAAT GAATACCATC 4680
 TGTTAACTCA GAAATTGGCT TAAAGTCAGA ACGTTCAACT TGTTCAGAA AATGTTGCAC 4740
 GTCGCGGCGA CAGGTCAGTT TTAGTAAGGT TTCAATCTCT CCATAAACAG GATGATCAAT 4800
 CAAGATATTT TGAACGCGAC CACCATTATC TACGATAGCA AGTAATTCTC GTCCAATTT 4860
 TTCAACTTCA TGCTTGACCT TAAATAATTT GTGATGATAA GTATTTGCAT TAGCATCTTT 4920
 ATAGATATAA CCACGATTGG TAGATAGAAT TGGAGATCCA TCAGCTCTTA AAATTGCAAT 4980
 ATCTTGAACA ATAACCTGTC GAGTGACATG AAAGTGCTCA 5020

(2) INFORMATION FOR SEQ ID NO: 143:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 4965 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 143:

AAAAAGTGGC AATCCATTGA TTGGCCACTT CATTTAGAGA ATTATCGTCT CGCCCTTGAA 60

967

GAAGAAGGTC GTGTAGTACT TGAGTTACTG CTATCGCTAG AACTACTACT TGAAGTCTG	120
GAGCTGGATG GAGTTGGTAG ACTCCCCACA ATACTAGACC AAGCATTCTG ATAATCCGCA	180
TCACTTCCGC CAATAGCAAA GCGATAACTT GTCGCTGGCG CTCCTGACTT ATTAGCCCAA	240
TAGCTGGTAA CAGTCGAACC TGTGACCTCT ACTTCTTTTC CTCAACAGA AACCTTCTCT	300
GGTTTTTGAC CTGTTGATTT CAAGACTTCC GATTTCACTA CACTAGGATC TAAAGCAAAG	360
CGCTCGTTCC CCCAAATGCT TGGGAAGCT TGCTGAATCG CATTACCAG ATGAGCCATG	420
TAATTAGAGT TATTAGAATA ACCTGCTCTA CGTGACAATG AATGATTATC ATCATGCCCA	480
ATCCAGCCAC CTAGGGTTAA TCTAGGTGTC GAAAGCATGA GCCACATATT TTCGTCTTGG	540
TTGGTTGTAC CAGTCTTCCC AATCCAATCT GCATTAGCCA GAGTAGGATT TAAAGAAGTC	600
AGTTAGACT TGAAGGTTGT TGTCACACGA GAGGATAGAA CTCTCGTAG CAATCCCTGC	660
ATAATCGTCG CAGTAGCTTT TGAATAGACT TGAACCGGTT TATCCTGATA CTCATACACC	720
ACTCTACCAT CTGCTGCTTC AATCTTTGAA ATCACATGCT TCTGATGATA AACTCCATTA	780
TTAGCTAAGG TCTGATAGCC ATTGGTATGC TGGGCAACTG TGAATCAAT ACCACCACCC	840
ATTGGCAAGC TCTCAATACC GFACTCAGGA ATCTCGTAAC CCATCTTTTC CATATAACCC	900
TTGACATCAA CACCCTTTTC ACGGAGCATA CGTAGGTCC AGTAAGCAGG GATATTCCAT	960
GAATAGTTCA GAGCTTCTCC CAAGGTCATC ATCTCTGTTT CCTTGCTATT AGCATAACATA	1020
ATCGGATTGC CATTAGCAAA GTTTGTTGGA TAGTTAGATA GAATCGTTTC ACTTCCCATC	1080
AAGCCCTGGT CAATAGCAAT ACCGTAGGCC AGCAAGGGCT TGGTAGTAGA AGCTGGCGAA	1140
CGTTTGGTAT CAAAGGCATG ATTATTTTGA TTTTCTTGAT AATTACGACC ACCTACAAAG	1200
CCTAGAATAG CACCTGTTTG GTTATCCATC AAGACATPCC CTAATCTTAC ACGACCTGTT	1260
CCATCGTCTA AAAGATAGCC ATAATCAGCA ACCGCACTTT GCATGGCAGA ATGAATTTTC	1320
TGATCTATGG TAGTAGTAAT CTTATAACCA CCATTTTCAA TTTCTTGGC TGCCAAATCT	1380
CGATAAAACT TCTGAGTTGC CTCATTTTTC AACTCCTTAG CGGAGACATT GTCTCTCTGA	1440
GCTAGATAGT CATAACATAC TTCTTGAGCT TCTGCCAAAG TTGTAAAGTA TAAATAGTCT	1500
CGTGAAATTC CTGTAACCGT GCCCGATGGT AAAAAGTCCT GTTTAAGGTC ATAATCCTTG	1560
TACTGAGAACT ACTCGTCTTT GCTTAATGCA CCTGTACGAT ACATACTGTA AAGAACTGCC	1620
TTAGCCCGTC TTAAGCCAAT TTCTAGGTCT TCATCACTCT TCAACTCCCC AGTATTTTCA	1680
TAAGGAGAGT AAGTAATGGG ACTCTGTGGA AGTCCTGCTA AAAATGCTGC TTGAGGAACA	1740
GTCAACTGAC TGGCATCTAC ACCGAAAATT CCCTCAGCTG CTGCGGAGC CCCTGCAATA	1800

TTCTGTCCCT	TATTATTTTCG	GCCAAAGGGA	GCCACATTGA	GATAGGTTCGT	TAAAATCTCA	1860
TCTTTATCA	TGGCGCGTTC	CAAGGCAAGA	GCATCCACAA	TCTCTGCCGC	CTTACGAGCC	1920
AAGGTCGGCG	CATCCCCAAC	CACCTGCTGT	TTAATTAGTT	GCTGGGTCAA	GGTTGAACCC	1980
CCACTAGAGG	AACCCAAACC	TACAAATTTT	CCCAAGGTCG	CACGAATCAC	CGCCTTGGGT	2040
ACTACACCCT	TATGTTCTTT	AAAGTGTTCA	TCTTCTGTCG	CAATGATAGC	CTTCTTCAGA	2100
TTTTCCGAAA	TTTGCTCAGA	TGAGATAGAA	GTGCGCAACA	AATCACTCTC	TATGGAAGCA	2160
ATCACCGTCC	CGTCCGAATA	GGTAATCTCT	GAAATAGAAG	AGATGTCCTT	GACCTGATTC	2220
ACCAATTCCT	CTGTCTGAGG	CACCCGAACC	TTGTCAAATA	AGGCCACTCC	GTATCCCAA	2280
GCAATCCCAG	CTCCCAACAT	TCCTCCTAGA	AAACCGAGTA	CAAAGAGTAA	GTAAATAAG	2340
GCTTTTATAC	TCAGTAAAAT	AGCTGGGAAA	ATGACTGACT	TATCTAAGGT	TTTAGATTTT	2400
TTGGTACTTG	AACCTTCTCT	GCCAGGTCTA	GCTGATTTT	TATTTTTTTG	TTTTTGCTGG	2460
AAAAATCCA	GCATTTTTTCG	TTTTAATCA	TTTAATTGAT	TTTGCATGGA	TTTCCTCACT	2520
TTATCTATTA	TACCACAAAA	GGGAAATTTT	CAATAAAATA	GCCACTTCT	TCCCTATCT	2580
GCTAGGCTAT	TGCCCAAGTT	TGTGATACAA	TAGGTAGAAA	CAATAATTTT	AAAAGGAGA	2640
AAAAACACAT	GCACATTTTT	GATGAGCTAA	AAGAGCGTGG	TTTGATATTT	CAAACGACTG	2700
ATGAAGAAGC	TTTGCCTAAA	GCCCTAGAAG	AAGGTCAAGT	TTCTTATTAT	ACTGGCTACG	2760
ATCCAATGC	TGACAGCCTT	CACCTAGGCC	ACCTTGTCGC	AATCTTGACA	AGTCGTCGCT	2820
TGCAACTAGC	AGGTACACAA	CCTTATGCGC	TCGTTGGCGG	TGCTACAGGT	CTCATCGGAG	2880
ATCCGTCCTT	CAAAGATGCT	GAACGTAGTC	TCCAAACAAA	AGACACAGTA	GATGGCTGGG	2940
TCAAGTCTAT	CCAAGGACAA	CTTCTCGTT	TTCTTGACTT	TGAAAATGGC	GAAAACAAGG	3000
CTGTCATGGT	CAACAACACT	GAAGGTTT	GCAGCATCAG	CTTCATGAC	TTCTCCGTG	3060
ATATTGGAAA	ATACTTCACG	GTCAACTACA	TGATGAGTAA	GGAATCTGTT	AAAAACGGGA	3120
TCGAAACAGG	AATTTCTTAC	ACTGAGTTCG	CTTACCAAAT	CATGCAAGGG	TATGACTTCT	3180
TCGTCTTAA	CCAAGACCAT	AATGTCACTC	TTCAAATCGG	TGGTTCTGAC	CAGTGGGGAA	3240
ATATGACAGC	TGGTACCGAA	TTGCTTCGTC	GTAAGGCCGA	CAAGACTGGT	CACGTTATCA	3300
CTGTTCCACT	AATCACAGAT	GCAACTGGTA	AGAAATTTGG	TAAATCAGAA	GGAAATGCCG	3360
TCTGGCTCAA	TCCCGAAAAG	ACTTCTCCAT	ACGAAATGTA	CCAATCTGG	ATGAACGTGA	3420
TGGACGCTGA	CGCTGTTCGC	TTCTTGAAAA	TCTTTACTTT	CTTGTCACTT	GATGAGATTG	3480
AAGATATTCG	TAAACAATTT	GAAGCAGCGC	CACACGAACG	CTTGGCTCAA	AAAGTCTTGG	3540
CTCGTGAAGT	TGTTACACTT	GTTACGGAG	AAGAAGCCTA	CAAAGAAGCA	CTTAACATCA	3600

969

CTGAGCAACT CTTTGCAGGA AACATCAAAA ACCTTTCCTGT CAAAGAGCTC AAACAAGGAC 3660
 TTCGTGGTGT GCCCAACTAC CAAGTACAGG CAGACGAAAA CAACAATATC GTGGAAGTGC 3720
 TCGTCTCATC TGGTATAGTT AACTCAAAAC GCCAAGCCCG TGAAGACGTC CAAAACGGAG 3780
 CCATCTACGT AAACGGCGAC CGCATCCAAG AGCTTGACTA TGTCTTGAGT GACGCTGATA 3840
 AGTTAGAGAA TGAAGTACT GTTATCCGTC GTGGGAAGAA AAAATACTTT GTATTGACTT 3900
 ACTAACTAT TCAACATTTA TCTATAAACA AAGGAGTTAA CCTCGAGAAA GGTAACCTCT 3960
 TTTTGCTGTT AATAACTCTC ATCTATCTAT TTTTAATAGA CAGGCTACGC AGGACAATGC 4020
 GCAAGGTTGT TAGATTATGT AAGATAGAGA GATTTGAAGG ACTGAACCAA TTAAATAAGC 4080
 CAAAGCCAAT CAAACTACTA TTTACGACAA CGGTATCCTG AATATTTTTC TTGATGAGTG 4140
 TTTGCAAAGA TGATGATAAC GAATCCAAC CTGGAAGAA ATCCAAACGA TTATCTAACA 4200
 ATAAGATATC ACTCATCTGC TTAGAAATAT CTGCACCTC ATTCATCACC ACACCGATAT 4260
 CTGATAGAGT TAAAGCCGCT GAGTCATPCA ATCCATCTCC AACCATCAA ATAGTGTGAC 4320
 CTGCTTCTG CAGTTTCTCT ACTAACTCAA ATTTCCCATC AGGTTTCAAG TCTGTATAGA 4380
 CCTGATCAA GGGCAAATCT TTGACTAATT CCTCTGTCTT AATCAAGGTG TCTCCTGTTG 4440
 CCAGAATCAA TTTTTTCCCC TGTGCCTTAA GTTTATCCAA GGCTGTTTTT GCTTCTTTTC 4500
 TCAAAGGAGT ATGAATGCAG AACATTCCAA TCAATTCATT TTGATAAGCC AAGAATAAGA 4560
 GATTGTAGTG ACTCTGTGAC TCTTCAATTA AAGCATTTTG TTCTGAACTG ATATGAATCT 4620
 GCTCATCTG CATCAAGACA TAATCCCAA TAAGAAGTGG TTGGCCATCT ATATGAGATT 4680
 TGATCCCCTT GCTTGCGATA TATTGGAGTT TCCCATGCAT TTCCTCATGT TCAATTCCTT 4740
 CTATCTCAGC TTGCTTGACG ATGGCATTAG CAATAGGATG ATAAATGTGT TCCTCAAGAC 4800
 AGGCACTGAT TCTGAGAATA TCTTCTCAC TATAGTCTCC AAAAGGTAAC ACCTTTTCAA 4860
 CTATAGGATA ACTAGTTGTG ATTGTTCCCTG TCTTATCAA CAAGAAAGTA TCAACTTCCA 4920
 GATATTTCTC CTGTTGTGG CCTCTGGCTG TCATCTCTGT GCTGG 4965

(2) INFORMATION FOR SEQ ID NO: 144:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 3232 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 144:

970

CAGGGGCGTA TTACGTGACA ATTCAATGTA GGCTGTCGCT ACTTGCGCCA AAACAAGGAT	60
TCGATAATGT CCGATGATAC TAACGATTAA ACCGAGCAGA AAGGATCCCA AAATTCCCCA	120
AACTGCAATA TGCAAGGTCA GAAAGAATGC CTTTTGATAT AGTGGTAGAT ATTGTTCAAC	180
AATGGATCAA TCCAAAAATA GAACCTCCCA TCTAGAAATA ATACAGTTAT TGTAGCACTT	240
AAAACTTCT TTGGATAATA TCTATTTTTT ATTGCCGTA TAAGGATTTT TATCATAGAC	300
ATAAAATTC TGAAATTTCC AAACAAAATA TTTTAAAAGT TTTGAAAAG AGTTAAGATA	360
TTTTTGTAAT ACACAAAGTA AACGCTTACT TATTAAGGAG GACATTTTAT GTCATACAAA	420
ACAAGCAATG CAGAAGGTCA TGTAGATTTC ATCAATACCT ATGATTTGGA GCCAATGGCG	480
CAACAAGTTA TTCCTAAAGC AGCATTTGGC TATATCGCTA GTGGGGCGGG AGATACTTTC	540
ACTTCTTCC AGTGATTTTA GCGTCAGGTT CTTTTAGTT TTTAAAGATT ATCCGTGAAT	600
TTCTTGCTTA TTTATGATAA AATGGGAGTG TCGCAAAAAA TGACTCATCG TATTCATTT	660
TGAGTAAAC TAGGAGGATC CCATGTCTAC AGAACATATG GAAGAACTAA ATGACCAGCA	720
GATCGTTTCG CGTGAAAAA TGGCTGCGCT CCGCGAACAA GGAATCGATC CTTTCGAAA	780
ACGTTTTGAA CGTACTGCAA ATTCACAAGA ATTAAAAGAT AAATATGCCA ACCTCGATAA	840
AGAACAATTA CACGATAAAA ACGAAACAGC TACTATCGCA GGACGCTTGA TAACCAAACG	900
TGGTAAAGGA AAAGTTGGTT TTGCCACCT TCAAGACCGC GAAGGCCAGA TTCAGATCTA	960
CGTTCGTAAG GATGCTGTCG GTGAAGAAA CTACGAAATC TTCAAAAAA CAGACCTTG	1020
TGACTTCTT GGTGTCGAAG GTGAAGTGAT GCGTACGGAT ATGGGAGAAC TCTCTATCAA	1080
GGCAACCCAC ATCACACACT TGTCTAAGGC TCTTCGTCCT CTTCTGAGA AATTCATGG	1140
TTTGACAGAC GTTGAACAA TTTACCGTAA ACGTTACCTT GACTTGATTT CTAATCGTGA	1200
AAGCTTTGAA CGCTTTGTCA CTCGTTCAA AATCATCTCT GAAATCCGTC GTTACCTTGA	1260
CCAAAAAGGA TTCCTTGAAG TGGAAACACC TGTTCTTCAT AATGAAGCCG GTGGTGTGTC	1320
TGCCCGTCCA TTTATACCC ACCACAATGC CAAAACATT GACATGGTGC TTCGTATCGC	1380
GACTGAGCTT CACTTAAAC GCCTTATCGT GGGTGGTATG GAACGTGTCT ATGAAATTGG	1440
CCGTATCTTC CGTAACGAAG GAATGGACGC TACTCATAAC CCTGAGTTCA CTTCTATCGA	1500
AGTTTACCAA GCTTATGCAG ACTTCCAAGA CATCATGGAC TTGACTGAAG GCATTATCCA	1560
ACACGCTGCT AAATCAGTCA AAGGTGATGG CCCAGTCAAC TACCAAGGTA CTGAAATCAA	1620
GATTAACGAA CCATTTAAGC GTGTTTATAT GGTGGATGCT ATCAGAGAAA TTTACTGGTGT	1680
CGATTTCTGG CAAGACATGA CTTTGAAGA AGCTAAAGCT ATCGCTGCTG AGAAGAAAGT	1740
TCCAGTTGAG AAACACTACA CTGAGGTTGG TCACATCATC AATGCCTTCT TTGAAGAGTT	1800

971

TGTTGAAGAA	ACTTTAATCC	AACCAACCTT	TGTCTATGGA	CATCCAGTAG	CTGTATCTCC	1860
ACTCGCTAAG	AAAAATCCTG	AAGACCAACG	CTTACTGAC	CGTTTCGAGC	TCTTTATCAT	1920
GACTAAGGAG	TACGGTAATG	CCTTACTGA	GTTGAACGAC	CCAATCGACC	AACTTAGCCG	1980
TTTTGAAGCC	CAAGCTAAAG	CAAAGAACT	TGGTGATGAT	GAAGCGACAG	GAATCGACTA	2040
TGACTACATT	GAAGCTCTTG	AATACGGTAT	GCCACCAACA	GGTGGTTTGG	GAATCGGTAT	2100
CGACCGTCTC	TGCATGCTCC	TCACTGATAC	AACAACATATC	CGTGATGTAT	TGCTCTTCCC	2160
AACAATGAAA	TAAATCTTA	TCCTCTGGGT	CTTATCAGAG	GATTTTTTGA	TTCAAAAAGA	2220
GACTGAATTT	AAGGAGAAAA	TGAAGTGTAG	TATATTGAAA	TTGAAATAGT	ACACTTTGAT	2280
TTCTAAGACA	TTGTTAGAAA	TTGGTTTAAA	TCCCTAAGC	AATTTGTGCA	TGTTTTATTT	2340
CATTTTACGA	TAGTACGCTG	AACTTTTCA	AAAAGTACTA	GAAATTGACT	TGGATTCCCC	2400
AATTGATTTG	TTCAGATTCA	CTATAAATA	AAAATTAATA	AGTGGGATAG	GAAGTTAGCG	2460
TCAACTAGGA	TAGTATCTTG	CTTAAACAGT	ATATATGGGA	TTGATATAAG	TCCATAGGTC	2520
CTATTAGAGG	ATGTTCTGGT	GTCTTATCA	CTTGTTTTTT	ATAGTATTAG	TAGATAGAAT	2580
CAGCAAATAA	AAACCCAAAT	CATTCATACC	TCTCTCAACT	AGATGTAACT	TACAAAACCC	2640
CTGACCTCAT	GAGCCACTTT	CTTCCTCCTC	ATGAGGTCAG	TTTTACTTTC	TGCTGTTCCA	2700
GTATCGTTTT	TCCTCGCTAG	ATTCCTCAA	AAGGGCAGAC	TCCTCCCTTG	GTGCGTCACA	2760
CGATTTTTTC	ATCTCGACTG	TTCTTTAATG	CATCATTAAC	GACGCTTTTC	TTCTAGGTGG	2820
TTCATAAGGA	ACAGGAAGAT	TCAGGTTGAC	TTTTCTAATC	CTAGAATAAA	GTGCTGAAAA	2880
CAATTCGGAA	TAGGCATAGA	GACTAGACAA	TTTGAGGAGC	TGCTTGCGTC	CTGTTCGAAC	2940
ACATTTTCCC	ACCACGTGAA	GAAAAAGATG	GCGGAAGCGT	TTGATTGTTA	AAGTTTGAA	3000
GTCACCTCCA	GCTAGATGTT	TGAGAAAAAG	ATAGAGATTG	TAGGCGATAC	AGCTCATCAT	3060
CATACGAACT	TCGTTTTTGA	TTAAGGTTGA	ACTATCCGTT	TTATCGCCAA	AAAATCCCTC	3120
CTTCATCTCC	TTGATGAAAT	TCTCGGCTTG	ACCACGTCCA	CGATAAAGCT	GAAACTGGTC	3180
TTGGCTTGTT	CCACTCGTCA	TATTTGTAAC	GAGAGAAATA	ACATCGTAGA	AC	3232

(2) INFORMATION FOR SEQ ID NO: 145:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 10711 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

972

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 145:

CCGGAGAAAA	TGATGAAAAG	TTCAAAACTA	TTTGCCCTTG	CGGGCGTGAC	ATTATTGGCG	60
GCGACTACTT	TAGCTGCATG	CTCTGGATCA	GGTTCAAGCA	CTAAAGGTGA	GAAGACATTC	120
TCATACATTT	ATGAGACAGA	CCCTGATAAC	CTCAACTATF	TGACAACCTGC	TAAGGCTGCG	180
ACACAAATAT	TACCAGTAAC	GTGGTTGATG	GTTTGCTAGA	AAATGATCGC	TACGGGAACT	240
TTGTGCCGTC	TATGGCTGAG	GATTGGTCTG	TATCCAAGGA	TGGATTGACT	TACACTTATA	300
CTATCCGTAA	GGATGCAAAA	TGGTATACTT	CTGAAGGTGA	AGAATACGCG	GCAGTCAAAG	360
CTCAAGACTT	TGTAACAGGA	TTAAAATATG	CTGCTGATAA	AAAATCAGAT	GCTCTTTACC	420
TTGTTCAGA	ATCAATCAAA	GGTTGGATG	CCTATGTAAA	AGGGGAAATC	AAAGATTTCT	480
CACAAGTAGG	AATTAAGGCT	CTGGATGAAC	AGACAGTTCA	GTACACTTTG	AACAAACCAG	540
AAAGCTTCTG	GAATTCTAAG	ACAACCATGG	GTGTGCTTGC	GCCAGTTAAT	GAAGAGTTTT	600
TGAATTCAAA	AGGAGATGAT	TTTGCCAAAG	CTACGGATCC	AAGTAGTCTC	TTGTATAACG	660
GTCCTTATTT	GTTGAAATCC	ATTGTGACCA	AATCCTCTGT	TGAATTTGCG	AAAAATCCGA	720
ACTACTGGGA	TAAGGACAAT	GTGCATGTTG	ACAAAGTTAA	ATTGTCAATC	TGGGATGGTC	780
AAGATACCAG	CAAACCTGCA	GAAAACTTTA	AAGATGGTAG	CCTTACAGCA	GCTCGTCTCT	840
ATCCAACAAG	TGCAAGTTTC	GCAGAACTTG	AGAAGAGTAT	GAAGGACAAT	ATTGTCTATA	900
CTCAACAAGA	CTCTATTACG	TATCTAGTTG	GTACAAATAT	TGACCGTCAG	TCCTATAAAT	960
ACACATCTAA	GACCAGCGAC	GAACAAAAGG	CATCGACTAA	AAAGGCTCTC	TAAACAAGG	1020
ATTTCCGTCA	GGCTATTGCC	TTTGGATTTG	ACCGTACAGC	CTATGCCTCT	CAGTTGAATG	1080
GACAAACTGG	AGCAAGTAAA	ATCTTGCGTA	ATCTCTTTGT	GCCACCAACA	TTTGTTCAG	1140
CAGATGGTAA	AAACTTTGGC	GATATGGTCA	AAGAGAAATF	GGTCACTTAT	GGGGATGAAT	1200
GGAAGGATGT	TAATCTTGCA	GATTCTCAGG	ATGGTCTTTA	CAATCCAGAA	AAAGCCAAGG	1260
CTGAATTTGC	TAAAGCTAAA	TCAGCCTTAC	AAGCAGAAGG	AGTCCAATTC	CCAATTCATF	1320
TGGATATGCC	AGTTGACCAA	ACAGCAACTA	CAAAAGTTCA	GCGCGTCCAA	TCTATGAAAC	1380
AATCCTTGGA	AGCAACTTTA	GGAGCTGATA	ATGTCAATTAT	TGATATTCAA	CAACTACAAA	1440
AAGACGAAGT	AAACAATATT	ACATATTTTG	CTGAAAATGC	TGCTGGCGAA	GACTGGGATT	1500
TATCAGATAA	TGTCGGTTGG	GGTCCAGACT	TTGCCGATCC	ATCAACCTAC	CTTGATATTA	1560
TCAAACCTTC	TGTAGGAGAA	AGTACTAAAA	CATATTTAGG	GTTTGACTION	GGGGAAGATA	1620
ATGTAGCTGC	TAAAAAAGTA	GGTCTATATG	ACTACGAAAA	ATTGGTTACT	GAGGCTGGTG	1680
ATGAGACTAC	AGATGTTGCT	AAACGCTATG	ATAAATACGC	TGCAGCCCAA	GCTTGTTTGA	1740

CAGATAGTGC	TTTGATTAT	CCAAC TACAT	CTCGTACAGG	GCGTCCAATC	TTGTCTAAGA	1800
TGGTACCATT	TACAATACCA	TTTGCATTGT	CAGGAAATAA	AGGTACAAGT	GAACCACTCT	1860
TGTATAAATA	CTTGGAACCT	CAAGACAAGG	CAGTCACTGT	AGATGAATAC	CAAAAAGCTC	1920
AGGAAAAATG	GATGAAAGAA	AAAGAAGAGT	CTAATAAAAA	GGCTCAAGAA	GATCTCGCAA	1980
AACATGTGAA	ATAACTGTTG	CAAAATATAA	GAAAGGATTT	AGTATTTCCC	TTGAATGCTG	2040
AATCCTTTTT	TACATTTGTA	AAGAAAGATT	CTAAAATGTA	CGGACCCCCA	AAAGTTGGAG	2100
CCTCTTTTTG	TCAGAATAGA	GAAAATTTTT	GTTAATTTTA	CTTGTTCCT	ATTGCTTCT	2160
CAGTATTAT	TTGTATAT	AAAAGTATAA	TTATTTTTTA	TTTATCAGAG	TTAAGCATTG	2220
CACTTCAGA	GGAAGGAGTA	TTTTTTAAAA	AGAAAATGTA	AACGTTTGCT	CAAAAATGAA	2280
AGGATTTAGA	AGTTTATGAA	TAAAGGATTA	TTTGAAAAAC	GTTGTAAATA	TAGTATTCGG	2340
AAATTTTCAT	TAGGTGTTGC	TTCTGTTATG	ATTGGAGCTG	CATTCTTTGG	GACAAGTCCG	2400
GTTCTTGCA	ATAGCGTGCA	GTCTGGTTCC	ACGCGAACT	TACCAGCTGA	TTTAGCTACT	2460
GCTCTGCAA	CAGCAAAGA	GAATGATGGG	CGTGATTTTG	AAGCGCCTAA	GGTGGGAGAA	2520
GACCAAGGTT	CTCCAGAAGT	TACAGATGGA	CCTAAGACAG	AAGAAGAACT	ATTAGCACTT	2580
GAAAAAGAAA	AACCGGCTGA	AGAAAAACCA	AAAGAGGATA	AACCTGCAGC	TGCTAAACCT	2640
GAACACCTA	AGACGGTAAC	CCCTGAATGG	CAAACGGTAG	CGAATAAAGA	GCAACAGGGA	2700
ACAGTACTA	TCCGAGAAGA	AAAAGGTGTC	CGTACAACC	AACTATCCTC	AACTGCTCAA	2760
AATGATAACG	CAGGCAAACC	AGCCCTGTTT	GAAAAGAAGG	GCTTGACCGT	TGATGCCAAT	2820
GGAAATGCAA	CTGTTGATTT	AACCTTCAA	GATGATTCCTG	AAAAGGGCAA	ATCACGCTTT	2880
GGTGCTTTTT	TGAAATTTAA	AGATACCAAG	AATAATGTTT	TTGTCGGTTA	TGACAAGGAT	2940
GGCTGGTTCT	GGGAGTATAA	ATCTCCAACA	ACTAGCACTT	GGTATAGAGG	TAGTCGTGTT	3000
GCTGCTCCTG	AAACAGGATC	AACAAACCGT	CTCTCTATCA	CTCTCAAGTC	AGACGGTCAG	3060
CTAAATGCCA	GCAATAATGA	TGTCAATCTC	TTTGACACAG	TGACTCTACC	AGCTGCGGTC	3120
AATGACCATC	TTAAAAATGA	GAAGAAGATT	CTTCTCAAGG	CGGGCTCTTA	TGACGATGAG	3180
CGAACAGTTG	TTAGCGTTAA	AACGGATAAC	CAAGAGGGGG	TAAAAACAGA	GGATACCCCT	3240
GCTGAAAAAG	AAACAGGTCC	TGAAGTTGAT	GATAGCAAGG	TGACTTATGA	CACGATTCAG	3300
TCTAAGGTCC	TCAAAGCAGT	GATTGACCAA	GCCTTCCCTC	GTGTCAAGGA	ATACAGCTTG	3360
AACGGGCATA	CTTTGCCAGG	ACAGGTGCAA	CAGTTCAACC	AAGTCTTTAT	CAATAACCAC	3420
CGAATCACCC	CTGAAGTCAC	TTATAAGAAA	ATCAATGAGA	CAACAGCAGA	GTACTTGATG	3480

974

AAGCTTCGCG	ATGATGCTCA	CTTAATCAAT	GCGGAAATGA	CAGTACGCTT	GCAAGTTGTA	3540
GACAATCAAT	TGCACTTTGA	TGTGACTAAG	ATTGTCAACC	ACAATCAAGT	CACTCCAGGT	3600
CAAAAGATTG	ATGACGAAAG	CAAACACTT	TCTTCTATTA	GTTTCCTCGG	CAATGCTTTA	3660
GTCTCTGTTT	CTAGTAATCA	AACTGGTGCT	AAGTTTGATG	GGGCAACCAT	GTCAAACAAT	3720
ACGCATGTCA	GCGGAGATGA	TCATATCGAT	GTAACCAATC	CAATGAAGGA	TTTGGCTAAG	3780
GGTTACATGT	ATGGATTTGT	TTCTACAGAT	AAGCTTGCTG	CTGGTGTTFG	GAGTAACTCT	3840
CAAAACAGCT	ATGGTGGTGG	TTCGAATGAC	TGGACTCGTT	TGACAGCTTA	TAAAGAAACA	3900
GTCGGAAATG	CCAACATATGT	AGGAATCCAC	AGCTCTGAAT	GGCAATGGGA	AAAAGCTTAT	3960
AAGGGCATTG	TTTTCCCAGA	ATACACGAAG	GAACTTCCAA	GTGCTAAGGT	TGTTATCACT	4020
GAAGATGCCA	ATGCAGACAA	GAACGTTGAT	TGGCAAGATG	GTGCCATTGC	TTATCGTAGC	4080
ATTATGAACA	ATCCTCAAGG	TTGGGAAAAA	GTTAAGGATA	TCACAGCTTA	CCGTATCGCG	4140
ATGAACTTTG	GTTCTCAAGC	ACAAAACCCA	TTCCTTATGA	CCTTGGATGG	TATCAAGAAA	4200
ATCAATCTCC	ATACAGATGG	TCTTGGGCAA	GGTGTCTCFC	TTAAAGGATA	TGGTAGCGAA	4260
GGCCATGACT	CTGGTCACTT	GAACATGCT	GATATTGGTA	AGCGTATCGG	TGGTGTGCGA	4320
GACTTCAAGA	CCCTAATTGA	GAAGGCTAAG	AAATATGGAG	CTCATCTAGG	TATCCACGTT	4380
AACGCTTCAG	AAACTTATCC	TGAGTCTAAA	TACTTCAATG	AAAAAATCT	CCGTAAGAAT	4440
CCAGATGGAA	GCTATAGCTA	TGGTTGGAAC	TGGCTAGATC	AAGGTATCAA	CATTGATGCT	4500
GCCTATGACC	TAGCTCATGG	TCGTTTGGCA	CGTTGGGAAG	ATTTGAAGAA	AAAACCTGGT	4560
GACGGTCTCG	ACTTTATCTA	TGTGGACGTT	TGGGGTAATG	GTCAATCAGG	TGATAACGGT	4620
GCCTGGGCTA	CCCACGTTCT	TGCTAAAGAA	ATTAACAAAC	AAGGCTGGCG	CTTTGCGATC	4680
GAGTGGGGCC	ATGGTGGTGA	GTACGACTCT	ACCTTCCATC	ACTGGGCAGC	TGACTTGACC	4740
TACGGTGGCT	ACACCAATAA	AGGTATCAAC	AGTGCCATCA	CCCGCTTTAT	CCGTAACCAC	4800
CAAAAAGATG	CTTGGGTAGG	GGACTACAGA	AGTTATGGTG	GTGCAGCCAA	CTATCCACTG	4860
CTAGGTGGCT	ACAGCATGAA	AGACTTTGAA	GGCTGGCAGG	GAAGAAGTGA	CTACAATGGC	4920
TATGTAACCA	ACTTATTTGC	CCATGACGTC	ATGACTAAGT	ACTTCCAACA	CTTCACTGTA	4980
AGTAAATGGG	AAAATGGTAC	ACCGGTGACT	ATGACCGATA	ACGGTAGCAC	CTATAAATGG	5040
ACTCCAGAAA	TGCGAGTGGA	ATTGGTAGAT	GCTGACAATA	ATAAAGTAGT	TGTAACCTCGT	5100
AAGTCAAATG	ATGTCAATAG	TCCACAATAT	CGCGAACGTA	CAGTAACGCT	CAACGGACGT	5160
GTCATCCAAG	ATGGTTCAGC	TTACTTGACT	CCTTGGAACT	GGGATGCAAA	TGGTAAGAAA	5220
CTTCTACTG	ATAAGGAAAA	GATGTACTAC	TTCAATACGC	AGGCCGGTGC	AACAACCTGG	5280

975

ACCCTTCCAA	GCGATTGGGC	AAAGAGCAAG	GTTTACCTTT	ACAAGCTAAC	TGACCAAGGT	5340
AAGACAGAAG	AGCAAGAACT	AACTGTAAAA	GATGGTAAAA	TTACCCTAGA	TCTTCTAGCA	5400
AATCAACCAT	ACGTTCTCTA	TCGTTTCGAAA	CAAACATAATC	CTGAAATGTC	ATGGAGTGAA	5460
GGCATGCACA	TCTATGACCA	AGGATTTAAT	AGCGGTACCT	TGAAACATG	GACCATTTCA	5520
GGCGATGCTT	CTAAGGCAGA	AATTGTCAAG	TCTCAAGGGG	CAAACGATAT	GCTTCGTATT	5580
CAAGGAAACA	AAGAAAAAGT	TAGTCTCACT	CAGAAATTAA	CTGGCTTGAA	ACCAAATACC	5640
AAGTATGCCG	TTTATGTTGG	TGTAGATAAC	CGTAGTAATG	CCAAGGCAAG	TATCACTGTG	5700
AATACTGGTG	AAAAAGAAGT	GACTACTTAT	ACCAATAAGT	CTCTCGCGCT	CAACTATGTT	5760
AAGGCCTACG	CCCACAATAC	ACGTCGTGAC	AATGCTACAG	TTGACGATAC	AAGTTACTTC	5820
CAAAACATGT	ACGCCTTCTT	TACAAC TGGA	GCGGACGTCT	CAAATGTTAC	TCTGACATTG	5880
AGTCGTGAAG	CTGGTGATCA	AGCAACTTAC	TTTGATGAAA	TTTCGTACCTT	TGAAAACAAT	5940
TCAAGCATGT	ACGGAGACAA	GCATGATACA	GGTAAAGGCA	CCTTCAAGCA	AGACTTTGAA	6000
AATGTTGCTC	AGGGTATCTT	CCCATTTGTA	GTGGGTGGTG	TCGAAGGTGT	TGAAGATAAC	6060
CGCACTCACT	TGTCTGAAAA	ACACAATCCA	TATACACAAC	GTGGTTGGAA	TGGTAAGAAA	6120
GTCGATGATG	TTATCGAAGG	AAATTGGTCA	CTCAAGACAA	ATGGACTAGT	GAGCCGTCTG	6180
AACTTGTTTT	ACCAAACCAT	CCCACAAAAC	TTCCGTTTTG	AAGCAGGTAA	GACCTACCGT	6240
GTAACCTTTG	AATACGAAGC	AGGATCAGAC	AATACCTATG	CTTTTGTAGT	CGGTAAGGGA	6300
GAATTCAGT	CAGGTCGTCG	TGGTACTCAA	GCAAGCAACT	TGGAAATGCA	TGAATTGCCA	6360
AATACTTGGA	CAGATTCTAA	GAAAGCCAAG	AAGGCAACCT	TCCTTGTGAC	AGGTGCAGAA	6420
ACAGGCGATA	CTTGGGTAGG	TATCTACTCA	ACTGGAATG	CAAGTAATAC	TCGTGGTGAT	6480
TCTGGTGGAA	ATGCCAACTT	CCGTGGTTAT	AACGACTTCA	TGATGGATAA	TCTTCAAATC	6540
GAAGAAATTA	CCCTAACAGG	TAAGATGTTG	ACAGAAAATG	CTCTGAAGAA	CTACTTGCCA	6600
ACGGTTGCCA	TGACTAACTA	CACCAAAGAG	TCTATGGATG	CTTTGAAAGA	GGCGGTCTTT	6660
AACCTCAGTC	AGGCCGATGA	TGATATCAGT	GTGGAAGAAG	CGCGTGCAGA	GATTGCCAAG	6720
ATTGAAGCTT	TGAAGAATGC	TTTGGTTCAG	AAGAAGACGG	CTTTGGTAGC	AGATGACTTT	6780
GCAAGTCTTA	CAGCTCCTGC	TCAGGCTCAA	GAAGGTCTTG	CAAATGCCTT	TGATGGCAAT	6840
GTGTCTAGTC	TATGGCATA	ATCTTGAAT	GGTGGAGATG	TAGGCAAGCC	TGCAACTATG	6900
GTCTTGAAAG	AACCAACTGA	AATCACAGGA	CTTCGCTATG	TTCCCGTGG	ATCAGGTTCA	6960
AATGGTAACT	TGCGAGATGT	GAAACTTGTT	GTGACAGATG	AGTCTGGCAA	GGAGCATACC	7020

TTTACTGCAA CTGATTGGCC AAATAACAAC AAACCAAAAG ATATTGACTT TGGTAAGACA 7080
 ATCAAGGCTA AGAAAATTGT CCTTACTGGT ACCAAGACAT ACGGAGATGG TGGAGATAAA 7140
 TACCAATCTG CAGCGGAACT TATCTTTACT CGTCCACAGG TAGCAGAAAC ACCTCTTGAC 7200
 TTGTCAGGCT ATGAAGCAGC TTTGGTTAAG GCTCAGAAAT TAACAGACAA AGACAATCAA 7260
 GAGGAAGTAG CTAGCGTTCA GGCAAGCATG AAATATGCGA CGGATAACCA TCTCTTGACG 7320
 GAAAGAATGG TGAATACTT TGCAGATTAT CTCAACCAAT TAAAAGATTC TGCTACGAAA 7380
 CCAGATGCTC CAACTGTAGA GAAACCTGAG TTTAAACTTA GATCTTTAGC TTCCGAGCAA 7440
 GGTAAGACGC CAGATTATAA GCAAGAAATA GCTAGACCAG AAACACCTGA ACAAATCTTG 7500
 CCAGCAACAG GTGAGAGTCA ATCTGACACA GCCCTCATCC TAGCAAGTGT TAGTCTAGCC 7560
 CTATCTGCTC TCTTTGTAGT AAAAACGAAG AAAGACTAGT ATTTAGTAAA ACCTCTTAAC 7620
 AAGATTACGG AAGCAGTCTC TATCTTTTCC AATGAGGTTT ATAGTACAGA AAAAGCCTGA 7680
 GAAGATGTCT TCTCAGGCTT TTGTTAAGCA CATAAATACA ATAGTGCTAT GACAAAATCA 7740
 CCCAGAAAAA TCTGGGTGAT AAATGTTATG GTTGTGCTGG TTGAGGATTC TGATTTTGT 7800
 GATCAGGGGT TGTATTTGAT TGTTGCGTAT TATTGTTAGG ATTGGTAGTC GTACTATTAT 7860
 TTGTGCTTGG AGTGGTTGAG CTAGACTGTG AAGTTGAACT ATCTGATGAT GAGCTTGAAC 7920
 TTTCAAGTTGA TGGGGGTTGT TGTGGAGCAG GTGAGTTCCA CGTAGAACGA GCACCATTTT 7980
 TAAATACGAA TTCTCCATTT CTGTAGAGCC CCTCTGGTAT ATTCCAATCT TCTGGATTGC 8040
 TTCTTCAGA CAGGTAGGTC ATCATAGAGC GGTAAACTTT GGCAGCGACC GTAAGGCCAT 8100
 TGCCTACAAG TGGTGTGAGA CGGTTAGAAT AGCCTGTCCA TACAGCCATT GAATATTTAC 8160
 GCGTATAGCC AGCAAATAGT TCATCAGGTG CTACAAATG AGAGGTCTTG ATGTGGTTTT 8220
 CAATTTCTC GTCTGTATAG TTAGAGGTTT CTGTTTTACC AGCCTGAGGG AGCCAAGCAA 8280
 GATAGGCAT TCGTCCAGTT CCATAAGTCA AGACTGTTTT CATCATGTCG GTCATCATAT 8340
 AGGCTGTCGT TTCCTTCATG GCACGAGTTC CGACATTAGA GAACTTTTT TCACTCCCAT 8400
 CACTAAAGAC GACTTTATGG ATATACATTG GTTTATAGTA AGTTCCACCA TTTGCAAAGG 8460
 CAGCGTAAGC AGCAGCCATC TTTTCACTAC TTGCTCCATA TTTTGTGCT GATTCGGTTG 8520
 TGTACTTGA AATGGCATT T GAGTAGTAA TACTTGGGTA GTCGATTCCT AGACCATTTA 8580
 GGAAAGTCTT GGC CGGTTG AGTCCGACCT TGTTTAGAGT TTCCACGGCT GGGACGTTTC 8640
 GCGATTGTG CAGGGCGTAT TGCAAGGTGA TGTGCCAAA GTAGCCCCTA TCCCAGTTAT 8700
 AAACAGGAGT ATTTGTCCCA GGGTAGTTAT AGGGCTCATC GTGAACGATA GTAGCAGTTG 8760
 AATCGTAGAC ACCGTACTCC AAGGCAGGAG CATAGTCTGT GATCGGTTTC ATAGTTGATC 8820

977

CCCAGTCGCG	GTTTGTTCCT	ACTGCTTGGT	TAATTCGGAA	GGAAACATTA	CTTGACTGAT	8880
GGCGTGCTCC	TAGCTGGGCA	ATGACTTTAC	CGTTAGAAAC	ATCAACAATG	GTAGAAGCGA	8940
CTTGCAATTC	ATCGTCTGGA	TAGGCAACGT	ATTCTGTCTGT	ATTGTAAATA	TCCCACAGAT	9000
GTTTTTGAGC	TTCTTGCTCT	ACATTTGTGT	AGACATCCAT	CCCAGTTGTG	AGTAGGTTAT	9060
AGCCTGTTTC	TTCTTCAACT	TGATTGATGA	CTTCCTTGAG	GTAATTATCC	ATGTAAGCAG	9120
GGTAATTACT	TGCTGATTTG	AGACTTTGTA	GTCCATCAGT	AATTGGTGTA	TTGACTGCTT	9180
TCTCATACTG	TTCAGCAGAG	ATGTAGCCTT	GATTTTTTCAT	TTCAGATAAG	ACCAAGTTTC	9240
GGCGGTCTTG	GGCTGCTTCT	GGATGTGAAT	AGGGGTCATA	TTGGTTTGGT	GCCTGAGGCA	9300
TTCCAGCCAG	CAAGGCTAAC	TGAGGTAAAC	TTAAATTATP	GAGGTCTTTA	CCATAGTAGT	9360
TTTGAGCTGC	TGCTGCATP	CCATAGTTCC	CATTAGACAT	GTAGACCTTA	TTTATATAGT	9420
AGGTCAAGAT	TTCTTGCTTG	GTTGCTTTTT	GTTCTAAC TG	AATCGCTAAC	CAAGCTTCCT	9480
GAGCCTTACG	AGAAATAGTC	TGGTCGGAAG	TCGAAGTTGA	AAAGTAAGTC	AACTTAATCA	9540
ACTGTTGGGT	GAGAGTTGAT	CCACCTGGA	GGGAATTGCT	TTGCAGATTG	CGCAAGAAAG	9600
CTCCCAGGAT	ACGGATGGTA	TCAATCCCCC	TGTGGTCGAA	GAAGCGATGG	TCTTCGATAG	9660
AAACGATTGC	CTTAACCAA	TCTGTGGGAA	TATCATTAGC	TTGGGCATTG	ACGCGGCGTT	9720
CAGAACCCAA	GTCAGCAATG	AGTTGATTTT	TATGTGCGTA	GATTTTACTA	GAAGTTGTTG	9780
CAACTAGTTT	ACTCTCGGAT	AGGCTAGGAG	CCTTGCTAAC	GTAGTAGAAA	AAAACCTCCTC	9840
CGCCTAAGAC	AATGGCTGCG	ATAACCAAGC	TTAAGAAGCT	AATGCTCAGA	TACTTGATTA	9900
GGCGCAGAAT	CGTTGGTTTG	TTTATCTTGT	TTTACCACCT	AATAAATGTT	CTTTGATAAC	9960
ATTGAGATAA	GGAATTTGAG	GGAAGGCACC	AGCCTTGATT	TCATATCCAT	ATTCTCGAAT	10020
ATATTCAAGT	GGCATTGATT	TTTGTCCCTT	ATCTTGATGA	TAGAAGCGAA	TCAAATCGAA	10080
TGCCGGCAAT	AAGTAGGTTT	CTTGCTGAGA	AGAAAAGTGA	AGAAGGACAA	AGCAGATTCC	10140
TTGTTGGGCA	AGGACTTGTT	CCATATGCTG	AATCTGATGT	GGATGAAAAT	TTTTTCATCGG	10200
AATCGCACGT	TTTTGTTTTG	TTTCCTTGAC	TTCAAAGTCG	ATGTAATATC	CATTATAAAC	10260
GCCAGAATAG	TCCGTCGTTG	AAGCTTGTCG	AAAATAGGCT	TCAACAATCT	TGGCACGACT	10320
TCGTTGTGGA	TAGTCCACTT	GTACGATTTG	AATAGGAGTT	GGTTTCTTAT	GTATAACAGC	10380
CAAGCCCTGA	GACAAATAGT	AGTCGTTGGT	AGCATTGATC	ATCTTTTCAA	AGGGTACCGA	10440
GCTCGAATTC	GTAATCATGT	CATAGCTGTT	TCCTGTGTGA	AATFGTTATC	CGCTCACAAT	10500
TCCACACAAC	ATACGAGCCG	GAAGCATAAA	GTGTAAGCC	TGGGGTGCCT	AATGAGTGAG	10560

978

CTAACTCACA TTAATTGCGT TCGCTCACT GCCCGCTTTC CAGTCGGGAA ACCTGTCTGT 10620
 CCAGCTGCAT TAATGAATCG GCCAACGCGC GGGGAGAGGC GGTTCGCGTA TTGGGCGCTC 10680
 TTCCGCTTCC TCGCTCACTG ACTCGCTGCG C 10711

(2) INFORMATION FOR SEQ ID NO: 146:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 11887 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 146:

TACATTCATT CCATCGGCTA CTCCATAATA CTTAGATAAA ACCATAGCTG AAGTCGAATA 60
 CGGATACTGT AAAGTATTAT CAATTTTAAT CAAATCATCA TTACCGATAA TACTTCTGAT 120
 TGCTTTTGGT AGTATGAACC ATACGTTGGT GAAATCTCAG ATAATGAAGA ATCATTAGAC 180
 TCTGGACCTT TTTCTAGTGT CTCACTTACC TCATATCTT CACCCTTACT AGAAATAACA 240
 CTCAAAGCAG ATACTGTCGA TAACTGGCTA GCCAATAAAG TACTCGCAAT AATTGAAATA 300
 CCCAATTTTT TATAAACAGT TTTCTTCATT ATTGTATCCT CCTAATGTAA TTATAGCGTA 360
 CTATCTCTAAA TTTCTTAATC TACTATAGAA TCAAGAAATC TACCACCTTC TTTAAATACC 420
 CTCCATTATC ACATAAACAG GTAAACTTTT CAATTAATGA CTGCGCTTTT CAATCAGCT 480
 AGAGTACTT GCTTGCTTCT TTGATACTAA GTTCAGCCAT TCTTTCCTTG TTTTCTCAA 540
 TAAAGCATGT TACCCAAGTG GGATTCGTTT TGGAGTAGTC TCGCAGAGTC CAGCCAATGG 600
 CTTTATTGAT AAAAAATCT GTTTGGTTCA AGTTATGAAG GAGAATCTTT TCCATTAATT 660
 GAGTATTGGT CTTCTCTTTT CTTAACAAC TGGTGGTCAAT AGCGACACGT CTCAGCCAGA 720
 TATTATCTGA TAGGCTCCAT TTTATACTCA ATGAAAATCA AAGAGCAAAC TAGGAAGCTA 780
 GCCGAGTTG CTCAAAACAC TGTTTGGAGG TTGCAGATAG AGCTGACGTG GTTTGAAGAG 840
 ATTTTCGAAG AGTATTAAGA TTATTTCTTC TAGTTCAGGG TGTTCATACA CCAAACCTCC 900
 TACTACTCGA TCTAGGATAT CTACCGTGTC CCACAAGGAT TTTGTCACGA CTAAGTCTC 960
 TAGCTTAGGC AAATCGGTTT CCTTTAGATA AGACTGCATT GCTTTCAAAT AGTTAGCAGC 1020
 CACATATTGG TATTTTCTAG GATCCTTTTC CCAGCAAGTG TCTGCAAAAT CCCAATCGAT 1080
 AATCTTTGTT TTTTTCGCTT CTGGAAAATA TTTTATAGAG TTTATTTCTT TCAGGCACCG 1140
 CAATACCTAG AAAAGAAAAT TGATGGCGCA TATAGGCTTC CATGGACCTT GCTTTTTTAG 1200
 AGTCTTTTGC TGCTTCTAGC TCCTCAAGTA AATCTGCTAA ACTCATCTAA AACTCTCTT 1260

GCCCCACCAA	ATGGTGCTGA	AAGGCATAGA	CAGCCGCCTG	GGTACGATCG	CTGACTTCAA	1320
GTTTGGCAAG	AATATTGGAC	ACGTGGGTCT	TGACCGTCTT	GAGAGAGATA	AAGAGGTCAT	1380
CTGCGATGCG	CTGATTTTCG	TAGCCCTTGG	CGATGAGTTG	GAGAACATCT	CGCTCACGCG	1440
CAGTCAATTC	TTCATGAAGT	TCCATATGAT	TGCGGTGGTA	TPCAACCTTC	TGCTAACCT	1500
CTTGCTCAAT	GGCCAGCTCG	CCAGCAGCTA	CCTTACTGAC	GGCATGAAGC	AATTCATCTG	1560
CACTAGAAGT	CTTGAGCATA	TAGCCTTTGG	CACCAGCATC	TAAGACTGGC	ATGATTTTTT	1620
CATTGTCCAA	ATAAGAGGTC	ACAATCAAAA	TCTTGGCTTC	AGGCCATPCT	TTAAGGATTG	1680
CTAAGGTCG	GTCAATCCCA	TTCATCTCAG	GCATGACAAT	ATCCATGACA	ATGACATCTG	1740
GACGCAGTTC	CAAGGCCAAG	TCAATCCCTT	GAGACCCGTT	GGACGCCTCA	CCCACAACCT	1800
CTACATCGTC	TTGGAGGTCA	AAGTAGCTTT	TCAAGCCCAA	TCGGACCATT	TCATGGTCAT	1860
CTACTAGTAA	AATTTTCATC	TTTACTCCTT	TATCATTCCT	TATCTAACAG	GGGAATACGG	1920
ATATCAACCG	CCAGCCCTTG	CTTGGGAGCT	GTCAAGAGTT	GAAGTGTCC	AGCCATATCT	1980
TCAACCCGCT	CCTTGATATT	TCGCAGTCCA	TAACTCAAGT	CGTCTAAGCT	CCCTAACTGG	2040
AAACCAATCC	CATTGTCCAC	CACCTTCAGT	TGCAATTCAA	CATCTGTCTG	ATAGAGGTAG	2100
ACATCTAGGC	AAGATGCCTG	GGCATGGCGG	AGGGTATTGC	TAATCAACTC	TTGCAGGATA	2160
CGGAAGATAT	GCTCCTCGAT	TTTCTTAGGC	AATTCGTCA	TATTCTGCTT	GAGACTAACC	2220
CTAAGATCAC	TCTTGTCCTC	AAGCTCTTTT	AAAAGAATTT	GAATCCCTTC	TATCAAGCTC	2280
TTCTGCTCCA	GTTCAACTGG	TCGCAAATGC	AAGAGCAAAA	CCCGCAAATC	CTTCTGGGCT	2340
GTTTCTAAAA	TAGCTGTGAC	ACTCTGCAAC	TGGGTCTGCA	TCTTTTCTCT	ATCCAATTTT	2400
AAAGCCTGCT	GACTGATACC	CGATAAAAATC	ATGTGGGCCG	CAAACAACCTC	CTGACTGACT	2460
GTATCGTGCA	AATCCCGAGC	AATFCGCTTC	CGTTCCTTCT	CGATGATTTT	CTCTTCCTGA	2520
GCAAGGCTCT	GATTTTCAGC	TTTTTGAAGA	GCCTCTGTCA	AAAGGTTAAG	TTTACCTGAT	2580
AAGGACTTGA	AACTGGCATC	CAAATCTGGA	TCTGCAACCT	GAACCACTTC	TTGCCCTGCT	2640
AATAAACGCT	TGAGATTAGC	CTGCATTTTT	CTTAGAGAAA	GCTCTTCGAT	CCCTCGCCAA	2700
AACAGGGCTA	AGAGACAGGT	CATGGACATG	CTGAAAACCA	ACAATAAAAA	GACAAATTTT	2760
TCTGTTTTTT	CGACATCGTG	CAAAAAGATA	GACCAGTCAA	AATCAAGTAT	TTCCAGCAAG	2820
CTGTGGGAGA	AAAAAAGAC	AAATAGGAAG	GAGGTGAGAG	CAATAATGAC	ATAGGCTTGT	2880
TTTTTCATCC	TCTAACCACT	TCCACATCAC	CAATCATAGT	GGTCAAGAAA	ATCTTGACAC	2940
TCTTGTTACT	CTTGAGATAG	TCTTTGTTTT	CTTGATGATA	GTGTTCAATTG	CGGAGGGCTC	3000

980

GCTTGGGCTG	GTTGAAAAA	ATCAAATCCC	CATAGAGACA	GTTAACGCTG	AGACTGACTT	3060
CCACATCTAC	AGGTACGATG	ATTTTGGTCG	TTCCCTACCAT	CTTCTGAGG	ATAATGACAT	3120
TGTCATGATT	GGTTAAGATG	ACCCTCTCCA	GATGAATAGT	GTCCTTGCCC	ATGAAGCGAA	3180
AGAGATTGAT	ATCATCGAAT	TGGCAAGTCT	GGTAGCTTGA	AAAATGATGA	AGATTTCCAA	3240
ACCAACGATT	TTTCTCCTTC	TTAACCGTCA	CGACCTCTTC	AAAAACCAA	TTGGTCTGCT	3300
CTTTTCCTG	GTTTCATCATC	GGGTAAAGAA	GAAAGAGGCT	ATAGATAACC	GCAACAAAA	3360
TAGCTAGAAT	CACAAAAGGA	TTGAGCATAA	CGATGAAAAA	GAAGAGAATG	GTTGCCGCTA	3420
CTAAAAGAAG	ATTATTTCCC	TCTTTACCAG	TGTAGTAGCG	AATCAAAAGC	AAAAAGAGGA	3480
ATAGTATCAG	CAGAAAACGC	GAAAAATGCT	CTGATACCAT	CAAATCAGA	GCTCCTGTCA	3540
GAAGACAGGC	TTCGATAAAT	AAAAAGATTT	TAAATTTTCT	CATAGGTTC	TCTCTCCCT	3600
TCTATTTTAT	CACAATCAA	AAAAGTCACC	TCAGTCTGAG	GATGGAAAA	AGGCGCTGGT	3660
TACGCCTTTT	TCATCTGATC	CTTTGCTTCT	TTTAATTTTC	CATAAAGAAG	ATAGTCTACT	3720
TTTTGTAGAT	CTGCTATGGT	GGCACAGTTA	AGGGAACACA	TAATCAAGCG	TAGATCTGCT	3780
TTCCAGCCTT	GGACAATGCC	AATCACTTCT	TCAACTGTGT	AGGTTTCAAC	CAATTCAGCA	3840
ACGGTTCGTG	ACAATCCCAC	AGCCTTAGCA	CCAAAAACCA	AGCACTTAAT	CATATCCAGC	3900
GGATTCGGAA	CCCCCTCCACT	AACCAAGAGT	TCGACCTTAT	CTTTCCATTC	TTGGGCATTG	3960
AGAAGGGCCT	GCATGGTAGA	CTGACCCCAT	TGATTGAGGT	AATCACGCTG	GCCACTACGA	4020
CGGTTTTFCGA	TATAGGCAAA	GCTGGTGCCA	CCACGACCCG	ATAGGTCCAC	TGTACGAACA	4080
CCGAATFCAT	AGGCTCTTTC	GATTGCTTTG	GCATCCATTC	CAAAGCCCAC	TTCCCTGAGG	4140
ACAATAGGAA	CGGGAATTTG	CTTGCTATAA	TCTGCTAGAT	GCGATTGCCA	GCTTCTAAAC	4200
TTCCTTTCTC	CCTCGGGCAT	GAGTAATTCC	TGCATGACAT	TGACATGCAC	TTGCAATAGA	4260
ACAGGATFCA	TCTCTTCTAC	AGTCTGAAGT	CCTAATCGA	CAGGCTTGTC	CAATCCAATA	4320
TTGGTTCCAA	GGAGGAGATT	GGGATGACTA	GACTTGACAG	AAAAAGAATC	ATCCGTGGA	4380
TTTTTGAGGG	CTGCGCTATA	AGAACCCGTT	ACAAATAAAA	TACCACAGGA	TTCCGCCACC	4440
TGAGCCAGCT	TTTGATTGAT	TTCTCTTCCC	TTATTACTTC	CACCAGTCAT	GGCATTGATA	4500
TAAAAAGGAA	AGTCCCCTT	TCGACCAGCA	AACTCTGTCG	AAAGATCGAT	TTCATCCAGA	4560
TTGTAAAGAG	GCAAGGAAGA	ATGAATCAGC	TCCACCTCAT	CAAAGCTATT	ATAGGAACTT	4620
TTCTGCTCAA	GGGCATAGAG	GATATGCTCG	TCCTTACGAT	TTGTCGTCAT	GTCCTATCCT	4680
TTCTTGATAT	AAGAGCTCAA	TCCCCAGATC	GGCCCAACGA	TTTTTTAAGG	TTTTGGTTGA	4740
TTGCGCATCA	AAACTCAGGG	CGATGCCACA	GTCACCACCA	CCAGCACCAC	TACTCTTGGC	4800

AACGGTCTGC	AAATCTTGAC	TGGCTTCTTT	CAACTGTCTA	AGCAAAGGCG	TGTAATATC	4860
TGTACTCAAG	CCTTCTAAAA	GCTTGCTGGC	TACTTCTACT	TGATCGATAA	TCTTTTCTGA	4920
TTTCCCTGT	TCCAAGGCTT	CTACCAGAGA	AGTCACCGTT	TCTTTTGAGG	AAGTTAAAAA	4980
ATTTTGATTG	ATATTTGCT	TGATTTGCTG	GACCATGTGA	CTCGATACAG	CCACTTCCTT	5040
GGTCCATCCC	ACTAAGAAAT	CACATTCTAA	AGTTGGTTTC	ACTTGTGAAA	TTGAAAAGCC	5100
CCAATCACGC	TCCAGAACTG	TCGCCAAGTT	TTCTTCTTCT	AACCAAGCAG	CCACCTTCTG	5160
GCGATCAAAT	GACTGGTAGA	GAACCAAATC	CTCTGCCACA	ATACAGGCAA	GGTCGCCCAT	5220
GGAACCATTG	TCTCCTCGCT	TAAGCAAGAC	AGCGCTAGTC	AGCTTGAACA	AGAGCTCCTG	5280
ATCAACAGAA	ACATCATAA	GAGCCAGTAA	AGCCTTGACA	ACCAAGACAA	CGACGCTGCC	5340
ACTAGAACCT	AGACCAAAT	TTTTCCCTTC	TCGTTCCATT	TTGCCACAGA	TTTCTAGAGA	5400
AAAAGTCTT	AAATTCTGAC	CACGAACAGC	GAGGAAGTCT	CCCATCAAAG	CAATCGTTTC	5460
TTGAATCAAG	CTATAGTCAG	GATTAGGCCT	TAAGTCCACT	GCGAAATCAA	ACATATCTGA	5520
ATAGATACGG	TAGTGTGAG	AAAAAGCAAT	CTCAGCCCTC	ATATAGATGG	GAATATCCTT	5580
TATCAAAGCT	AACTGCCCTG	GCTCTAAAAT	AGCATATTCA	CCTGCCCAAT	AGAGTTTTC	5640
GCAAGTTTTA	ACAGCAATCA	TCTTGACTCA	AATCCTTTGT	TTTTGACACA	ATCAAGCGAT	5700
AACGATGACC	GAAAATTTCT	GATAAATGCT	CCAAGTCTTT	CTCCTGACAG	AAGACCTTAA	5760
CATTGGGACC	AGCATCCATG	GTAAAGTAGC	AGGCCTCTCC	TTTCTCACGA	AGCTGGCGAA	5820
CAAAGGCCAT	AGCCTCATAA	GAGGCATCCG	TCAGATAAGA	AAAGGCTGGA	CTAGCAGTCT	5880
TTGTCTAGC	ATGCATAGCC	AGGGCATTTT	TCTCCGTTAA	TTCTCCAATC	TTGGCAAAT	5940
CATTTTCCTT	GAGATAAATC	AGCATATCCT	GATAGTCCCT	CTCAGACTGA	CGAACCCAGT	6000
CGTCGAAAGT	CGTCGAGGTT	TCCACACAAA	GTTTCATCCC	GTCACGGCTA	GAGATTGGTT	6060
TTTTCTTGTC	CTCTAGCACC	AACATAATCA	TAGCTAGTTT	CAAGTCTGTC	TCTACAGGGT	6120
AAATTTCTCC	ACTATCCTTA	TCCCAGGCTC	CTAGTGGTCC	ATAAAAATC	CGAGAAGAAG	6180
AACCTGAGGC	AAATTGGCT	TCCTGTGCCA	ACTGACTTCT	ATCCAATCCA	AGCTTGAAAT	6240
AAGCATTACA	AGCCTTGACC	AGGGCGGACA	AACCACTAGA	ACTTGAGGAC	AGACCCGCTG	6300
CCGTAGGCAT	ATTGTTTTGA	GTATCGATAC	GGACAAAGCC	CTCACCAGCT	GGACGATAAC	6360
GGTCAATAAT	CTTACTCATC	TTGGCATGCT	CGACCTCAT	TTGTAGCTGA	CCATTGATGT	6420
AAAATTCGTC	AGCTGTTACA	TTGGCTGGTA	AAGGCGACAA	GGTCGTCTCT	GTATACATAT	6480
TTTCCAAAGT	TAGAGAAATA	CTGCTAGTAG	CAGGCACCAT	CTCTTTTCT	TTTTTCTTTC	6540

CCCAATATTT	GATAATAGCA	ATATTTGCGT	AGGAACGTAC	TGTTACAGGC	TCTCTATCCA	6600
TGTCTGAACA	GCTCCTTTCT	CTTCTAATCT	TTCTGCTAGT	TCTTGTGCGT	GTGTCAAATT	6660
GGTTACCAAG	GCTATGATAC	AACCTCCTAG	CCCACCACCG	CTCATCTTGG	CACCCAGAGC	6720
ACCATGGCTA	AGAGTCGTTT	CAACCAAAAA	GTCTGCCTCA	GGGCTACTGA	CTCCAATTTT	6780
TTTFAAATGT	AAATGCGCTT	GACTGAGGAT	TTGTCCCAGT	CCTTCAGCAT	CTTTTGTGA	6840
AATCGCAACT	TCTGCTTGCT	GGGTTAATTC	TCCCAAGGCA	TGCAAAAACG	GTAGGGCATC	6900
CTTGCCCTTA	TTTTGAACCA	CTTGGATGGC	TTCACGAGTA	TGACCATAAA	CACCCGTATC	6960
GGCAATCACC	AAATAGGCGG	ATAAATCCAT	CTCAAGTTCT	GTAATCCTA	CGTTCCTGAT	7020
AAAGCGAATA	GGTTGGTCAC	TAAGACAGGT	CTTAGCATCC	AAACCACTAG	GATTCATATG	7080
GGCAATCATT	TCAGCTCGAT	TGACCAAGAT	TTCTAGTACA	TCATGAGGCA	GATCAGCCTG	7140
ATAGTAGTCA	AATACTGCAC	GAATGGCCGC	TATGCTGATA	GCCGCTGACG	AACCCATCCC	7200
CCGTTTCTCA	GGGATAGCCG	AGTCAATCTC	ACAACGAATG	CAGGCTTCTG	TGATATTCAA	7260
ATACTCCAGT	GAGGCATAAA	CCGCCATGGA	CAAGGTATCC	TCCTCATAAA	GGCGCCAAGG	7320
ACTCTCTGCA	GGAACTACCT	TACAGGTCAC	CTCCACCTCC	AAAAGAGGCA	GGGAAATGGC	7380
AGGATAACCG	TAAACGACCG	CATGTTCCCC	TATTAAAAAT	ATCTTACTAT	GTGCCTGACC	7440
GACACCAACT	TTTTTTGTCA	TTTTTTCCTT	TTACTAGACG	AAAAAACGTC	TTATTTTTCA	7500
TACAAGTATT	AATCTTTTCC	TATCTATTTT	ATTATATTTT	CACAAAAAAA	GCGATTGTTT	7560
CCATTCACAA	TCGCTTCTTT	CATTATTGAA	CCCATTGCCC	ATTATAGTTG	ACAGAATAGC	7620
CATCTACGGT	CGTATTCACT	GCCAAGGCAC	CTGAGCGCTA	TAAGCGTAGT	ACCATCTGCC	7680
ATTGACCTGG	AACCAACCTG	TCGTCATAGA	ACGACGAAAG	AAACTCCATA	CCATTAAGTA	7740
AAGAGGAAAG	TCGTGAGGGA	GCATGCGCCA	TTGACAACCT	GTTTTAGTGA	CGTACAAAGT	7800
CTCATTAACA	AGTACTCGTT	TCGGCCATTT	ATAGGTGCGG	TGTTTGAGGA	AATAGGGTTC	7860
AATCTTCGCC	CATCTTGTAT	CGTTTAAATC	AGTATCATAT	GCTTTGCGTA	TCATAACTCT	7920
AGCTTAACAT	TTTTTTGTGA	ATACAGGTTC	TAAATAATCG	ACCACGAAAA	TTTCTTAAGT	7980
GGAAAACGCC	TTATGAAGTA	TGCTACGGGA	AAGTTATGCA	CTTAATTTGA	CAATTCAAGA	8040
TGTAAAAATA	TATACTATAG	TAGATTGAAA	CTAGAATAGT	ACACCTCTAC	TTCTAAAAATA	8100
TTGTTAGAAA	TCGATTTGAC	TGTCCTGATC	GATTTATCCT	GTATTATCT	CATTTTACTA	8160
TAATATTTGA	TAAGTTATCC	TAAAAGTATT	ATTATGTTGT	TGTGTTATAG	ATTGATTGAA	8220
TCTAACTAAA	GGATCCTATT	CAATTACTAG	AACTATCACA	TACTCAAGGT	CAGCTCACAG	8280
ATGAGCAACT	ATTTTGGTTA	CAATGTCTAC	TAAATTTAAG	TCAAACAAAT	AATTTAGTCA	8340

983

AAATTAAAA	AATAGAGGAA	CATAAATATG	ATTACAAAAC	AGAATGTAAT	AGTGTCTAC	8400
AATTTTTACT	AGATAAACT	GTAAATCTG	AAGGAAGGAT	CACTTCTTCA	ACAGAATTTG	8460
GAAATTTTCGT	AAGTAATTTA	TCATTCCAAC	ACGGAATAGC	TGGACTACTG	TTTCTCTTAA	8520
ATAAATTGTA	CCCCCAGAA	CTGGATTCTA	AAATACTCTC	TATCATCAAG	AAGGCAGTGA	8580
CAATTAGAAC	GACACACACA	TATGAATATC	AATACTCACT	GCTATTTGGT	GATGCAGGCT	8640
ATCTATGGTT	ACTCCTACAT	TTATTTTCTA	TCAGTAAAAA	TCAATACTAT	CTACAATTAG	8700
CAAACGTCAC	CGCTAAAAAA	TTAATAGAGA	ATTATGATAC	TCTAGAGGAA	ATAGACTTTG	8760
CATTGGGAAA	ATCTGGTGTC	CTATTATCAT	TAATAAAATA	CTATCAATTT	ACCAATGACA	8820
ATACTCTTAA	AATTTTCATC	CACAATAGTA	TAGGGGAAAT	TTATCATTAT	TTCTTACAAA	8880
GAGATACAGC	CAAAGAAAGC	ATTTTAGACT	ATAGCTTTGC	TCATGGATAT	TGTGGAATTTG	8940
CATATGCTTT	ATTTGCCTAT	TCTAAAGTCT	TAGAACCCTC	TATGTTTTAT	AATGATCTCC	9000
ATACATTCCA	TACTGAATTA	AAAAAATAT	TAGAAAAAGT	TACTTCTAAT	ACTGAAAAAT	9060
TAGGAAATTT	ACAACCTTCT	TGGTGCAAAG	GAATTTCCGG	AATAATCTTA	TATCTTTGTA	9120
TGTACGATTG	TGACGGAAAC	AAAGATATTA	TTAGTAAATA	TCAAGAATTT	GTTTTTAACC	9180
ATCATCTAAA	AATGATGACA	GGATATTGCC	ACGGAATAAC	TAGCTTACTA	CAAACCACTG	9240
TCTACAATCA	AAACAAATTA	CTGATGAAAA	AAATCCAACA	GGTAATTTTA	GCATGTTCTG	9300
AACGAGATGA	TCACGGTTTA	CTGATGTTTC	AAGGAGATAG	TGGTAAAGCA	GATTTGTTTG	9360
ACTTCGGAAT	AGGAAGCATG	GGGTATATTT	GTGTCTATTA	AATAATAAAT	TCCCATTCGA	9420
TGTGCAGACA	TAAGGAGAAA	AGTATGAAAT	TATTTTGGAC	AAACAACATA	TATAGACAGT	9480
TGCTGCTAAA	CAGCTGTTTT	TCATCATTCG	GCGACAGTAT	TTTCTACCTC	GCCATTATCA	9540
ATTATGTGGC	TCAGTACAAT	TTTCGCTCCG	TAGCGATTTT	ACTGATTTCC	ATTTTCAGAGA	9600
TGGTTCCCTT	ACTATCGCAA	CTCTTTCTCG	GGATTCTAGG	AGATTTTCAA	GAAAAATAGAG	9660
TCAAACACGC	ACTCTGGATT	GCCAAAATCA	AAATCCTGCT	CTACGCTATT	TTGACAGTAT	9720
TTCTCGTCTT	GTCGCCCTTT	TCATTAGTTT	CAGTCATTAT	GATTGTCATC	ATCAACCTCA	9780
TCTCTGACAC	CTTGAGCTAC	CTGTCTGCCT	ACATGATGAA	CGCCCTCTAC	ATCAGTGTAA	9840
TTAAGGACGA	CCTGCATGAT	GCCATGGGGT	TCAGGCAGTC	TCTGATGAGG	GTTGTCCGTA	9900
TTGTCCGCAA	TCTGGCTGGC	GCATTCCTTA	TCAATGTTAT	AAGTATTCAA	ACTATTTCCC	9960
TTATCAACAC	TCTGACTTTT	GTCATTGCCT	TTTTGGGCCT	GTATGTTATT	CGACATACCT	10020
TGTATGAGGT	TGAAAAAAGA	ATTGAAATGT	CACATACAGC	ACTGAGTTTT	AAGAAATATT	10080

984

TTCAACATCT	TAAACAGTCG	CTGGCTGTGC	TCCTGAGGTT	AAAAGATACC	GTCATACTAC	10140
TGTTTCTGAC	GACCAGTATG	ATTGCCATCT	TGGATGTGTC	CCCTCGGCTG	ATTGCCCTCC	10200
GCTTCATCCA	ACAGACACTA	GCACAACCTGA	GCATTGGGCA	ACTCCTCGCC	CTGCTCTCCA	10260
TCATCATGTC	TTGTGGAGCT	ATCCTTGCCA	ATATGACCAG	CAGTAATCTA	TTTAAAAATA	10320
TCCGTTTCAC	GCACCTCTTG	GTTTTCTGTG	AGATTTCCCT	ATTGACTCTA	ATAACTAGTA	10380
TCCTTTGTCA	AGCCTATATC	GTAATTTTCA	TGACCAGTTT	CATCAGTTCT	ACGATTATCG	10440
GCATTCTCAG	CCCTCGCCTA	CAAGCAGCTG	TCTTTGCCCA	TATCCCCAGT	GACAAGATGG	10500
GGACGGTGGG	CTCTGCTCTG	AGCACAGTGG	ACATTCTCGC	CCCGTCCCTG	CTCTCCCTAT	10560
TAGCCCTATC	CATAGCATCG	GGCGTTTCGG	TGCAGTTAGC	ATTGATATTT	TTGTATCTTA	10620
TTTTAATGTC	TCTTATCTTT	TGTCAATGGT	TAGTCAAGTT	CAACACTCAT	AACTAACGAA	10680
AAAGCATGTG	TAGATTTTAC	ATGCTTTTAA	TCTCCCAAT	CGTCAGGTCA	AGTACAACAA	10740
AGTCACTTCT	TTGATTAAGC	GAGTGTCTA	ATATAATTAT	AAGCGCCCTG	TCATTACCGA	10800
ACCCATTCGC	CATTATAGTT	GACAGAATAG	CCATCTACGG	TCGTATTCAC	TGCCAAAGCA	10860
CCTGAGCTAT	AAGCATAGTA	CCAGTTGCCA	TTGACCTGGA	ACCAACCTGT	CTTCATGTCT	10920
CCATTACCTG	CATTTAGGTA	GTACCAAGTT	GAACCATCTT	GATACCAACC	AGTTGCCATA	10980
GCTCCTGATG	AACGGAGATA	GTACCATTTG	TTCCCAAGGT	TTTGCCAACC	TGTTTTCATA	11040
TCGCCATTTG	GGTGGTCTAA	ATAATACCAA	GTGGTACCTT	CCTGATACCA	GCCAGTGGCC	11100
ATTGCTCCTG	AGGAACGGAG	GTAGTACCAC	TTATTACCTA	GATATTGCCA	ACCTGTTTGC	11160
ATAATACCAG	TTGTTGGATC	TAGGTAGTAC	CAAGTCGAAT	CATCGTTTAT	CCACCCCGCA	11220
CGTCTTTCAC	CACCAAGGTA	GTTTTCTCCA	TTAATTTCCG	TCTTAGCTAG	ATAATACCAG	11280
TTAGACTGAT	CATAAAGCCA	ACCTGTCTCT	AAAGAATGAT	TTTGATTAAA	GTAATAGTTC	11340
GTATAATAAC	GCTTCTCTTC	TTTATCTTCT	GAATCTTCAC	GTTTTTCCCC	GTACTTTCTT	11400
CCAACACTGT	CTTTAGTTTT	AATCTCTAAT	GTTTTCCAAC	CAACAAACTC	TTGTAGCACT	11460
CCATTTTTAT	CGAAGTAGTA	CCACTCTGAC	TTTGAAAAAC	CTTCTAATCT	GATACCATTT	11520
GGGTAAGGAC	CAATTGTA	ACCTTTAGAT	GGAAACGGGA	TATATTGCCA	GCCGACAACC	11580
ATCTCTCCAG	ATAGAGAATC	AAAATAATAG	TACTTACCAT	CAATCACTCG	CCAGTAGGTT	11640
TCTTTGAGGT	CCCCCTTTTT	GTAGTAGGTT	CTTCCGTTTT	CTTGACAAA	CTGCCATCCT	11700
TCAGAATCAT	CTGCAAATAC	TGTACTGGTC	CCTAGCAAAC	CAAAGAAAAA	TACTGTCACT	11760
CCAACTTGCA	TAGTTTTTTT	CAAAATTTTC	ATCTATATAC	CCTCCAATAT	TAAATCCACT	11820
CACCAGATGA	GGCGAAATTA	TAAACTTTAC	CATCGATAGT	TTGGCTACCT	GTAACCATTG	11880

CTCCAGG

11887

(2) INFORMATION FOR SEQ ID NO: 147:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 11340 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 147:

CCGGTATGTT CTGGAATACT ACCAATCTAA GCTGGCTGTG CCCTACAGTT TTACAACCCCT	60
GTACGAATAC CTTAAGGAAT ATGACCGATT TTTCAGCTGG GTTTTGGAGT CTGGTATTTTC	120
AAACGCTGAT AAAATATCCG ATATTCCTTT ATCAGTTTTG GAAAATATGT CTAAGAAAAGA	180
CATGGAATCC TTTATCCTTT ATCTACGTGA ACGTCCCTTG CTGAATGCTA ATACAACAAA	240
ACAAGGTGTT TCACAGACAA CTATCAATCG AACCTTATCA GCACTTTCTA GTCTTTACAA	300
GTATCTAACC GAGGAGGTTG AAAACGATCA GGGGGAACCT TATTTCTATC GTAATGTAAT	360
GAAAAAAGTT TCCACCAAGA AAAAGAAAGA AACCCTTGCT GCCAGAGCTG AAAATATCAA	420
GCAAAAACCT TTTCTAGGTG ATGAAACAGA AGGTTTCTA ACTTATATCG ATCAAGAGCA	480
CCCACAACAG CTTTCAAATC GAGCTCTCTC ATCATTCAAC AAAAATAAAG AACGAGATTT	540
AGCCATTATT GCCCTTCTCT TGGCATCTGG TGTTTCGCTTA TCTGAAGCTG TTAATCTAGA	600
TCTAAGAGAT CTCAATCTAA AAATGATGGT TATTGATGTT ACTCGAAAAG GTTGCAAACG	660
TGACTCAGTC AATGTGCGTG CTTTTGCTAA ACCTTATTTA GAGAATTATC TGGCCATTCG	720
GAATCAACGC TATAAACCGG AAAAAACAGA TACAGCCCTT TTTTTAACTC TCTACAGAGG	780
TGTTCCCTAAT CGTATCGATG CTTCTAGCGT TGAGAAAATG GTTGCTAAAT ACTCAGAGGA	840
TTTTAAAGTG CGTGTAACAC CCCATAAACT GCGCCATACA CTAGCAACTA GGCTCTATGA	900
TGCGACTAAA TCACAAGTTT TAGTCAGTCA CCAACTAGGA CATGCTAGCA CACAAGTCAC	960
TGACCTCTAT ACCCATATTG TTAGTGATGA ACAAAGAAT GCTCTGGATA GTTTATGATT	1020
TTACGTATTT TAAATTATGT AAATAAATAT CAAAAAAGA AGTTGGCCAA CTTCTTTTGG	1080
ATTTATCCAA CTACCGCTTC AGCGATTTCT TCACGGCTAA TACCAGCGAA GTAGCGTGTG	1140
ATATCAATGG TTTTtagcgc CTTAAGAACA TCTTCGCGTT CGTATTTTAC CCCACGAAGG	1200
ACATCTTCTA CTGCAGCAAC GTCTTCAATA CCAAAGAAGT CACCATAAAT CTTGATGTCT	1260
TGGATTTTGG ATTCAGTAAC GTTAGCAAAG ACTTCAACCT TACCACTAGT GAATTTGATT	1320

986

CCACGACGGA	CGTTAAATTC	AGGTGATTTA	CCATAGTTCC	AGTCCCAAGT	TCCAAACTTA	1380
GTATCCTTGA	TGCGATTGAT	TTCGGCCAAT	TCTTCTTCTG	AAAAGACGTA	TTCAGTCATC	1440
TCTGGGTACT	CTTTTTTCAT	GTATTCCAAG	AGTAAATCAC	GGAATTTTTT	GACTGTGATT	1500
TTTTTTGGTA	ATTCATTGAT	AATATTGGTT	ACACGGGCAC	GGACGGATTT	CACACCTTTT	1560
GATTCAAATT	TATCTTTTGA	AACCTTAAGG	GCATTTGCGA	GGACTGACAA	ATCAACGTCA	1620
AAGAGCAAGC	AACCGTGGTG	CATGATACGG	CCGTTGATAT	AGGCTTGGGC	ATTGCCACAG	1680
AACTTCTTAC	CATCAATCTC	AAGGTCATTA	CGACCTGTGA	ACTCAGCTTT	AACCCCAAGT	1740
TGAGCCAGGG	TATTGATAAC	CGGAGTTGAG	AAGCTCTTGA	AGTCAAATGC	CTTATTTTCA	1800
TCTTCTTTGG	AGATGATCGT	GTAGTTGAGG	TTATTTAAAT	CGTGGTAAAC	AGCTCCACCA	1860
CCACTAATAC	GGCGAACTAC	CTCAATACCA	TTTTCGCGAA	CATAATCACG	GTTGATTTCT	1920
TCGATAGTGT	TCTGGTGACG	ACCAACAATG	ATAGATGGCT	TGTTAATCCA	AAGTAGGAAG	1980
ATTTGATCCT	CATCCAAAAG	GTGTTTAAAG	GCGTATTCTT	CCAAGGCAAT	ATTA AAAAGCA	2040
GTGTCATTTG	AATGATTGAT	AATGTATTTT	ATGATATCCC	TTTACTTTAT	ATGATAGAAA	2100
CTGGAAATAA	CCTTCCAGTC	TAATCTATCT	TCGTTTATTT	TTTTCTTAGG	TGAATGGATG	2160
GCCATTCCTA	GAACATCTGC	AAACGCTTCG	TACATCACTT	CAGAGTAAAGT	TGGGTGCCCG	2220
TGGATGGTCT	TCAGCATTTT	CTCAACAGTG	ATTTCCATTT	CGATGATGCT	TGATGCTTCG	2280
TTTATTAATT	CTGCGGCTGC	AGGACCAATA	ATGTGTACAC	CAAGGATTTT	TCCGTATTTT	2340
TTATCAGCGA	TAACTTTTAC	GAAACCTTGA	GCTGCGTCAG	ATGCAATAGC	ACGACCGTTA	2400
GCAGCAAAGT	TAAACTTACC	GATGGCAACA	TCGTATTTCT	CACGGGCTTG	TTCTTCTGTC	2460
AAACCTACTG	CTGCTACTTC	AGGGAGAGTG	TAGATGGCTG	CAGGAGTCAA	ATTCAATTTG	2520
GCAACTGCAT	GATTTCCCTT	AAGGGCATTT	TCAGCGGAAA	CTTCACCCAT	GCGGAAAGCT	2580
GCGTGAGCCA	ACATCTTAGT	ACCGTTGATG	TCACCTGGTG	CATAAATGCC	TGGAAGTCAA	2640
GTTTCCATGT	ATTCGTTGAC	CTTGATACAA	CCACGATCCA	ATTCAAACCT	AACCTCTCUA	2700
ATACCTTCAA	GGTCTGGCAT	ACGACCAATT	GAAAGAAGAG	CTTTGCTTGC	GATGATATCG	2760
TCTTTTCCTT	CAACCTTGAT	ACGAAGTTGA	CCATTTTCCT	CAATGATTTT	TTGCAGTTTA	2820
GTACCAGTCA	AGATGGTCAT	TCCTTTACGC	TCAAGAATCA	AGCGAAGGTT	CTTAGAAACT	2880
TCCACATCCA	TAGCTGGAAC	TATACGGTCC	ATCATTTTCA	TAACAGTCAC	TTTTGAACCA	2940
AATGTCATGA	AGGCCTGACC	GAGTTCGATA	CCGACAACCT	CACCACCGAT	GATAACAAGG	3000
CTTTCTGGCA	CTTCGTTTCA	TTCAAGAATG	TCATCACTAG	TCATGACAAG	TGGAGATTCC	3060
ATACCAGGGA	CGTTGATCTT	GTTGACTTTT	GAACCACCAG	CAAGAATGAT	TTTCTTGGTT	3120

TCAAGCAATT	CAGAACCATT	TACCAAGACG	TTCTTGCTCT	TAGTGATTGT	ACCAATTCCT	3180
TTATGAACAG	TAACTCCGTA	GCTACGAAGA	AGTCCTGCAA	CACCACCAAC	AAGAGTATTA	3240
ACAACCTTAG	ATTTAGTTTC	TAAAAGTTTT	TCCATATCAA	CAGTGAAGTT	AGGATTTTCA	3300
ATCACGATAC	CACGATTTGC	AGCATGACCG	ATATTTTCAA	TAATTTTCAAGC	GTTATGAAGG	3360
TAGGTCTTGG	TTGGAATACA	TCCACGGTTT	AAGCAGGTTT	CACCAAGTTC	AGATTTCTCA	3420
ACAAGGGCAA	CCTTACCGCC	GAATTGGGCA	GCTTTAATGG	CTGCAACATA	ACCAGCAGGA	3480
CCTCCACCAA	TCACAACGAT	ATCAAAAAGCA	TCATCGCTCT	TACCATCATC	GTTTGAGGTA	3540
CTTGCTACAG	GTACAGGGCT	AGCTTCTGGC	GATGCTGCTC	CAGCTGTTGG	GATGTTTTTC	3600
CTTCTTTCAC	CAAGGTAACC	GATAACTTCC	GTTACAGGGA	CAGTTTCACC	ATCTCCTTTG	3660
AGAATGGCAA	TCAAGTACCC	ATCTTCTTCG	GCTTCCAATT	CCATGCTGAC	TTTATCAGTC	3720
ATGATTTCCA	AAAGGATTTT	TCCTTCTTTT	ACAAATCTCT	CGACTTTTTT	ATTCCATTTG	3780
ACGATTTGTC	CTTCTGTCAT	ATCCACGCCG	GCTTTTGCCA	TAATTACTTC	TAAGGCCATG	3840
TCTTCCTTCC	TTTATCTATA	TCTTAAAAAT	GAATACTCTT	GCTCTTAAAT	TAACATTGAG	3900
ATTGGCGTTT	CAATCAACTC	TTTCAAGTCC	TTCATAAACT	TAGCACCAGC	CATACCATCT	3960
ACGACACGGT	GGTCAATGGT	TAATCCTAAA	CTCATGATTG	GGCGAATCAC	AATTTACCCA	4020
TTGACGACAA	CTGGCTTCTC	GATTGTCGAA	CTGACACCAA	GGATAGCTGA	GTTGGGTTGG	4080
TTAATAATCG	GACCAAAGGA	CTGAACACCA	AACATTCCCA	AATTACTGAT	TGTGAATGTT	4140
GAATTTTGTA	ACTCACTTGG	AGCCAATTTA	CCATCCAAGG	TACGGCCAAT	AACATCCTTA	4200
AAGGCTACAA	CCAGTTCTGA	AAGACTCATC	TTCTCAGCAT	TGTAAACAAC	AGGTGTCATC	4260
AATCCATTAT	CCATCCCAAC	TGCCATGGCA	AGATTGACAT	AGTTGTGAGT	GATAATAGTC	4320
TTGCCATCTT	CTGTCAATGA	AGCGTTGATG	TATGGGTGTT	TCATAAGAGT	CTTAACAACCT	4380
GCAAGCGAAA	GAAGTCTGT	TACAGTAGTC	TTCTTCCCAG	TTGCTTCCAT	GATTGGCTCA	4440
AGAACCTTCT	TACGAAGAGC	CAACATTTCA	GTCATATCAA	CTTTCATAGTT	GAGGGTGAAG	4500
GTTGGCGCAG	TCAAGTAAGA	TTCAACCATG	CGTTGGGCAA	TAACCTTACG	CATTGGTGTC	4560
ATTGGAATAC	GCTCGATTTT	ACCATATGGT	GTTACGTTAT	CAGGGACTTC	TTCCACTTTT	4620
TCAATCTGAG	CAGGAGATTT	GATGCTATCG	TTTTCGATAT	TTTCAGGAAG	CAGGGCCAAA	4680
ACATCCTTCT	TCATGATTTT	ACCACGATGA	CCGGTTCCTT	GGATTTCCCTG	CCAAGCAATG	4740
TTATGTTTCGA	GGGCAATTCG	TTTTGCAAGT	GGCGAAATGC	GAACCACGTT	TGTGTCCTTA	4800
TAAGTTTCCA	CGTCTTCTTT	GTGGACACGA	CCGTTTGCAC	CTGAGCCAGA	AACGTCGTAG	4860

AGGTTTATCC CTAATCATC CGCTAACTTT CTAGCTGCAG GAGTCGCTCT TAGCTTGTC 4920
 TCAGCCATGA CCTCTCCAAT TCTATTTATG ATACAAAGGG CGTCAAAAGC GACTGAAAAA 4980
 TAGGAAATCG ACGATGGCTT CGATGAAGCC AAGGAGATTT ATCTTTTTC CGATCTTTTA 5040
 GCCCGTGCCT TAATCTAAGA TATTAATGAC GAAGAGCTCT GCACCTAAAA GATACAAAGT 5100
 TTCTCGTCAG CTTTATTTTA TTTACATAAC TTATCTTATG TAACCCTATT CTTTGTATA 5160
 AGTTTTTCGG ATTCATCTT TGATACTTTC AACTGTTGGA ATCATGTCAT TTTCTAGGTT 5220
 TTGTGCATAA GGCATCGGCA CATCTTCTCC TGCACAACGG CGAATTGGTG CATCTAGATA 5280
 GTCAAATGCT TCTGATTCTG AAATAATAGC TGAAATTTCA CCGATATAGC CACTTGTTTT 5340
 GTGGGCATCG TTGACCAGAA CAACCTTACC AGTCTTCTTC ACTGAGTTTA TGATGATATC 5400
 CTTATCAAGC GGAACAAGGG TACGTGGGTC AACAAATTTCA ACTGAAATTC CTTCTTCTGC 5460
 TAATTTCTCA GCAGCTTGAA CCACACGGCG AAGCATTTTT CCATAAGTAA CAACTGTTAC 5520
 ATCCGTTCCCT TGGCGTTTGA TTTCACCAAC CCCAAGTGGG ATTTGTGTAGT CTGGATCAAC 5580
 TGGCACTTCC CCTTTTGGT TAAATCTGA CTTGTACTCA AGTATAATAA CTGGGTTGTT 5640
 ATCACGGATA GAAGACTTAA GCAGGCCTTT CATGTCCGCA GGTGTTCCAG GTGCCACAAC 5700
 CTTAAGTCCT GGAATGTGAG TAAACCAAGA CTCTAGAGAT TGTGAGTGCT GGGCGGCAGA 5760
 GCCAACTCCG TTACCAGCTG CACAACGAAC AGTCATTGGA ACCTGACCTT TACCACCAA 5820
 CATGTAACGT GTTTTAGCAG CTTGGTTGAC GATATTGTCC ATGGCAATAA CAGAGAAGTC 5880
 CATGAAGGTC ATATCGACGA TTGGACGAAG TCCTGTCTATG GCTGCTCCTG CTGCTGCTCC 5940
 AGAGATGGCA GCTTCAGAAA TCGGACAGTC ACGGACACGT TCTGGACCAA ATTCTTCAAG 6000
 CATTCCAACA GAAGTACCGA AGTCTCCTCC GAAGACACCG ACGTCTTCTC CCATCAAGAA 6060
 CACATTTTCA TCGCGACGCA TTTCCTCAGA CATAGCAAGG ATAATGGTGT CACGGAAGGA 6120
 CATTGTTTTT GTTCCATTT TATCTCTTTC TCCTTAGTCT GCGTAAATAT CTTCAAAGGC 6180
 TGATTCAAGC GGTGGGAATG GGCTTTCCTC TGCAAATTTA ACAGAAGCTT CTA CTGCTTC 6240
 CTTTACTTGC GCTTGGATTT CTTCCAATTC TTCGGCACTT GCAATGTTAT TTCAATAAG 6300
 GTAATTGCGG AGGTTTTTGA TTGGATCTTT TTGTTTCCAC AATTCCTT CTTACGCGT 6360
 ACGATATTTA CCAGGGTCAG ATGATGAGTG ACCGAGCCAG CGATAAGTTA CACTTTCAAT 6420
 CAAGACTGGA CCATTGCCAC TGCGAACATG GTCCACAGCT TTCTGAAATC CTTCATAGAC 6480
 ATCGATGACA TTGTTACCGT CTTTCATGAA CATTCAGGA ATTCCATAAG CGGCGCTACG 6540
 TTGATGGATA TGTTCTATAT TGGTCATTTT CTTGATATCC GCAGAGATAC CGTAACCGTT 6600
 GTTAATGCAA TAGAAAATGA CTGGCAGGTT CCAGATAGAA GCCATGTTCA CTGCTTCGTG 6660

GAAAACACCT TCATTGGTCG CACCATCTCC AAAGAAGCAG ACAACGATTT TACCGGTATT	6720
TTGCATTTGC TGA CTGACCGGAC AGCGATCCCC ATACCACCAC CTACGATACC	6780
ATTGGCACCA AGGTTCCAG CATCAAGGTC AGCGATATGC ATAGATCCAC CTTTCCCTTT	6840
ACAGGTTCCA GTGTATTTAC CAAGGATTTT AGCCATCATT CCGTTGAGGT CAATCCCTTT	6900
AGCAATAGCT TGCCCGTGTC CACGGTGGTT TGAGGTAATC AGATCATCTG GATTGAGAGC	6960
TAACATAGCC CCCACGTTAG CTGCCTCTTC ACCAACAGAA AAGTGCCTCA TTCCTGGCAC	7020
TTTCCCTTTC TTTACTAATT GTGCAATTTT TAAGTCCATG CGACGGATTT CTTCATCTT	7080
ACGGAACAT TCTAGCAAAA GATTTTTATC TAAAGTTGAC ATCTTCTTGC CTTTCTAACT	7140
TTCTTCTTAC CTTACTATTT TACCGCTTTT GGCAAATACT GTCAAAGTTT TTCTAAAAGA	7200
AATTTACAAA AATAAAAAAG AAAACCCCGT GAAAACAAGG GATTTTCTTG TCAAGAATAT	7260
TTTTTCACAA ACTTTTTAGC ATTTGGATTT TGCTAAAGAT TCAAATCTCT TCATAATCAC	7320
AGTTAAACGC CAACGGTAGA GCGCCCCGCT CACAATCAAA CTAATAATCA AGCCGATCCA	7380
GTAAGAATAA GCTCCAAAAT CTGTTAGGGA ATCAAATAGC GTAnCACAGG GATTGCTACG	7440
CCCCAATAAC CAAGCAAACC AAGGTA AAAA GGAATAACTG TATCCTTATA CCCCCGAAA	7500
ATTCCTGAA GCGGCGCCGC AAAGGTATCT GCTAACTGGA AGAAAAGACT ATAAGTTAAA	7560
AAACGCACTG TCAAATCGAT AAATTTTGGG TCGTTACCAT AAAGACTGGC CACATTTCCC	7620
CTAAAAATGT AAAGGAAGGT TAAGGTGAAG GCCGCAAAA TGAGGGCAGT CCATCTTCTT	7680
AGACCAATAT AGTTTTTCGC ATCATCAAAT CGCTTGGCTC CCACTTCATA GGAAACGACA	7740
ATAGCCATAG CCGATGAGAT ACTCATAGGA AAGGCGTACA TAAGACTTGA AAAGTTCATA	7800
GCTGACTGGT GACTAGCTAT AATCAAGGGC GAAAACCTAG CCATAATCAA GCCAACCCTT	7860
GAAAAGATAG CCACTTCCGC GAAGACAGTT CCCCCAATAG GCAGACCTAA ACGAACTCCT	7920
TCCTTAATTT TATCCATATT AAGTGAATTT CGTTTCTCAA GGTGTAAGGC TTTGAGCTTC	7980
TCCTGTTTAA ATAAAACCAG AACAGAAATC CCAAGCAAGA CCCAGTAGGC CAAGGATGTT	8040
CCTAAACCAG CACCAGCCCC TCCCAGTTCT GGAACACCAA AGGCACCGTA AATCAAGAGA	8100
TAGTTAAATC CGCTATTGAG AGGGAGTAAC AAAAGCATGA GTACATGGA CAGTTTGGTC	8160
AAGCCCAGCG AATCCAGCAA GGAACGAATG ACGCTAAAGA GCAACAAGGG GATAATCCCG	8220
ATAGATAAAA ACCAAAGATA GCGAACCCT ACTGCCGCTA CTGCTGCTTC TAACCCAATA	8280
TGATTCAAGA TTATTGGTGC CAAGAAAAGT ACCATCCCA GCAAGACCAC AGATAGGCC	8340
AAGGCCAAAT AAATAAATG GTAAAAATCA GACGCAACTT CTTCCTTTT GCCTCGACCA	8400

990

AGATGGTGAC	CAATGATAGG	CACCAAGGCT	GACACAATCC	CTGTTAGAAA	TGTAAAGAAA	8460
GGATTCCAGA	TACTGGTTGC	CATAGATACA	CCAGCCAAGT	CCATAGTGT	GTATTGACCT	8520
GTCATTGCAG	TATCAACAAA	AGAGGCAGAA	TAATTGGCAA	ATTGGTAGAT	CAGGATTGGG	8580
AAGAAAATTT	TAAAAATAA	TACTAACTTC	TCTCGTAAAC	ACTTTGTCTT	ATACATACTT	8640
CTCTTCTAT	TCTGATTTAT	CTAAACCAA	GAGTTTCAGA	CCATAGTTTT	TCAAACCTAG	8700
CGGAGGTTTA	TTAGATTTTG	AAGTAGTATG	CCAACACGCA	CATGTACGAC	AATAATAGCT	8760
TCTAACTAAA	CCTCCGTAT	CATATTGAAC	CGCATGGTCA	GCTTTTTCTT	TAGTTTCATA	8820
TTGAATTTTG	GAACGATTAG	CTGCGGGACA	GTAATTCCTA	CTATTAGATT	TCGCTTGTCT	8880
CTCCCTACGT	TTTCGAAAAT	AATTCATATT	CTAACTCCTA	TCAAGCTTGA	TAGACGATTT	8940
GTCCCTTACA	GATGGTATAT	TTAACCTGCC	CTTTTAAGGT	TTCACCGATG	AATGGTGAAT	9000
TAGCTGCTTT	GGAAGCAAAA	TGGGAGTCCA	CAAAGCGGTC	AGCCTTGGCA	TCAAAAATAG	9060
TGATATCTGC	TGGACCATTC	TCAGCCAAGT	AACCTGCTTC	AAAGTTGTAA	AGCTTGGCTG	9120
GGTTGTATGT	CATTTTTTCA	AGTAATTCCTA	TCAAGCTCAA	CTCACCAGCT	TCTACTAAAT	9180
AGGTCAAGCT	GAGAGACAGG	GATGTTTCTA	AGCCAGTCAT	ACCAGATGGC	GCTTTGGTAA	9240
TATCCTCAAC	ATTTTTTCA	TCTACATGAT	GAGGCGCGTG	GTCAGTCGCA	ATAACTGTGA	9300
TGACACCTGA	TTTGAGACCT	TCGATAACGG	CACGACGGTC	TGATTCCAAA	CGAAGCGGTG	9360
GATTCATCTT	AGCATTGCTA	CCTTGTGTTA	AAAGAAGTGC	TTCTGTCTTA	GAGAAATGCT	9420
GTGGCGCTAC	TTCTGCTGTG	ACTTCTGCAC	CTAACCCCTG	AGCAAACCTCC	ACTACTTTAA	9480
CACTTCTTTC	CTTAGACAAA	TGCTGGATGT	GAACATGGGC	TTTAGTTGCA	TAGGCAATCA	9540
TGACATCAG	CGCCATCATA	GCGTACTCAG	CCACCCAGT	AGCACCGCAG	ATATGGAAAT	9600
GTTCTCTAGC	AATATTTTCA	TTAAAGCCAA	GAACACCGTT	CAAACCTGGA	TCTTCTCAT	9660
GAAGGCTGAT	AAAGGTATTG	AGTTTTTTGG	CTTCCTCCAT	GGCTTCCCTG	ACAATCTTAC	9720
TGCTCTCAAG	CGGAATACCG	TCATCAGAGA	AACCAACCGC	ACCAGCTTCT	AAGAGTGCTT	9780
TAAAGTCAGT	CAAGTTTTTA	CCATTAAAGT	TTTTAGTAAT	GGTCGCAACT	GTCTTGACAT	9840
TAATCTTCTC	TTTGGCAGCT	GACTGGAGAA	CTGCTTGCAA	AGTCTCCACG	TCTGAAATGG	9900
TTGGACTGGT	ATTAGCCATC	ATGACGACAG	TAGTAAAACC	ACCTGCAGCG	GCTGCTAGGG	9960
CACCAGTATG	AATGTCTTCT	TTATGTGTTT	GACCAGGTTT	ACGGAAATGA	ACATGAATAT	10020
CGACCAAGCC	AGGAGCAACC	ACAAGACCAG	TAGCATCAAT	CGTTTCTGCT	CCTTCTCCG	10080
TGATCTCAGA	CGCAATTTTG	ATAATTTTCC	CATCTTGAAC	TAAGACATCA	CAAACCTGAT	10140
CCAAACCAGA	CTTGGGATCC	ATTACACGAC	CATTTTTGAT	TAGTAGCATC	TGCTTCTCC	10200

TTTATTCATA GAAATCAACT TGGGTATCCA ACAATTTATC CCCATCATAA ACAAACCTGG 10260
 CTGAAAAGAA GGGTTTATCC TCTAAAAGCC ACTCAACAAA GGTGTGGTCA CCTTCCCAAG 10320
 TCGGCTTGCT CAAAACCTCA TCATAGGGAA CCCATTCTAG CGTCCCCTCA TTGCAGTCAA 10380
 TCAAGTCGCC CTCAAACCTCC GTCACCTTAA AAACATAGGT GTACCAGTCT AAATCTGGTG 10440
 TAAATTCAGG AAAAGTGATG ACACCTTTTA GAACTGGCTT GGCTTTGAGC CCTGTTTCTT 10500
 CAAGGATTC ACGCGCCGCG CATTCTGGG GCGTCTCTCC TCTCTCTAGC TPACCACCCA 10560
 CACCAATCCA TTTCCCTTCA TGGACATCAT TGGGTTTCTT ATTACGATGG AGCATGAGCA 10620
 GTTCTTTCCC ATTATCAATG TAGCAAATCG TCGCTAACTG AGGCATATTT TCTCCTTATC 10680
 TAAGCCAATC GATTGGCTCT TGTCCGTGCT CTTTTAAGAA TGCATTGGCC TTGGAAAAGG 10740
 GCTTGGAACC CAAAATCCT CTATAAACCG ACAAAGGACT TGGATGGGCT GATTTCGATAA 10800
 TCAAGTGATG AGGATTGGTA ACTAATGCCT TCTTCTTACG TGCATAAGCT CCCAGAGTA 10860
 CAAAAACGAC TGGTCTATCT AGATGATTGA CCACCTGAAT CACAGCATCA GTAAAAGGCT 10920
 CCCAGATTG ACCAGCATGA CCATTGGCCT GTCCAGCAGG AACAGTCAA CAAGCATTAA 10980
 GAAGCAAGAC TCCTTGCTCA GCCCAAGCTG TCAAATCATG AGATTTCTTA ACTCCGATAT 11040
 CATCTGACAA TTCTTTCAAG ATATTTTGCA AGGATGGTGG AGCTGGGATA GAGTCAGGTA 11100
 CAGAAAAACT CAAGCCCTGC GCTTGACCTG GTCCGTGATA GGGGTCTTGC CCTAGAATTA 11160
 CCACCTTAAC TTCTTCAAGC AGTGTGTC AAGAGAGCCTG AAAAAACCTTT TCCTTGGGTG 11220
 GATAAATAAT CCCCTGAGAA TAGACCTGCT CCATAAATG ATTGATTTTC CCGAAATAAC 11280
 CCTCAGGTAA TTGCGCCTTA ATCAAAGCAT GCCAAGACGA GTGTTCCATA GCCGACTCGG 11340

(2) INFORMATION FOR SEQ ID NO: 148:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 12127 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 148:

AAAAAATAGA CTTGTTAGAC TATAAATGTA GTAAGCCTAC ACAAGAAAAA TACATAGAGA 60
 TAAAGGTGAT TATTATGAAA TTCAAAAAAA TGCTTACTCT TGCAGCCATT GGCTTATCAG 120
 GATTTGGGCT TGTTGCCTGT GGCAATCAGT CAGCTGCTTC CAAACAGTCA GCTTCAGGAA 180
 CGATTGAGGT GATTTACGA GAAAATGGCT CTGGGACACG GGGTGCCTTC ACAGAAATCA 240

CAGGGATTCT CAAAAAAGAC GGTGATAAAA AAATTGACAA CACTGCCAAA ACAGCTGTGA 300
 TTCAAAATAG TACAGAAGGT GTTCTCTCAG CAGTTCAAGG GAATGCTAAT GCTATCGGCT 360
 ACATCTCCTT GGGATCTTTA ACGAAATCTG TCAAGGCTTT AGAGATTGAT GGTGTCAAGG 420
 CTAGTCGAGA CACAGTTTTA GATGGTGAAT ACCCTCTTCA ACGTCCCTTC AACATTGTTT 480
 GGTCTTCTAA TCTTTCCAAG CTAGGTCAAG ATTTTATCAG CTTTATCCAC TCCAAACAAG 540
 GTCAACAAGT GGTCACAGAT AATAAATTTA TTGAAGCTAA AACCGAAACC ACGGAATATA 600
 CAAGCCAACA CTTATCAGGC AAGTTGTCTG TTGTAGGTTT CACTTCAGTA TCTTCTTTAA 660
 TGGAAAAATT AGCAGAAGCT TATAAAAAAG AAAATCCAGA AGTTACGATT GATATTACCT 720
 CTAATGGGTC TTCAGCAGGT ATTACCGCTG TTAAGGAGAA AACCGCTGAT ATTGGTATGG 780
 TTTCTAGGGA ATTAACCTCT GAAGAAGGTA AGAGTCTCAC CCATGATGCT ATTGCTTTAG 840
 ACGGTATTGC TGTTGTGGTC AATAATGACA ATAAGGCAAG CCAAGTCAGT ATGGCTGAAC 900
 TTGCAGACGT TTTTAGTGGC AAATTAACCA CCTGGGACAA GATTAAATAA AATGTTTGCT 960
 CCATAAATCT CTAAGAGAT GCAGACGTTT CATCGTACAA TAAGATAAAG AAGGCAAGTA 1020
 GGGAGGTGTC GTATCTCCCT TACTTTCTTC ACTAGAAAGG ACAAGATGTG ACAAACAAG 1080
 CCTTCAAAGA AGCAGTTTTT AGGGCAATTT TTTTCATGAG TGCAACAGTA GCTGTTGTAG 1140
 CTATTTTGCT AATCTGTTC TTTATTTTTA GTAATGGCTT ACCTTTCATA GCTAACTACG 1200
 GCTTTGCCCG TTTTTTATTA GGCAGTGATT GGTCCCAAC GAACATTCCG GCAAGCTATG 1260
 GTATTTTACC AATGATCGTT GGTCCCTTAT TAATTACCTT AGGAGCGATT GTGATTGGGG 1320
 TGCCAACAGG CATCTTGACA TCGGTGTTTA TGGTTTATTA TTGTCCAAAG CCCGTCTATG 1380
 GCTTCTTAAA ATCAGCTATC AACTTGATGG CAGCCATTCC ATCTATGTGTT TATGGTTTTT 1440
 TCGGCCCTACA ATTATTGGTG CCTTGGATTA GAAGCTTTTT AGGAAATGGC ATGAGTGTCC 1500
 TAACCGCTTC GTTACTATTA GGAATAATGA TTTTGCCAAC CATTATCAGT TTGTCAGAAT 1560
 CTGCTATCCG AACAGTCCC AAAACGTATT ATTCTGGTAG CTTGGCTCTA GGAGCTAGTC 1620
 ATGAACGGAG TATTTTTAGT GTCATCTTGC CAGCTGCGAG ATCTGGTATT TTATCAGCAG 1680
 TTATTTTAGG AATCGGTGCG GCAGTAGGTG AAACCATGGC AGTTATTTTG GTGGCAGGCA 1740
 ACCAGCCGAT TATCCAAGT GGACTCTTTT CAGGAACCAG AACCTTAACA ACCAATATTG 1800
 TTCTGGAAAT GGCTTACGCA TCAGGTGAGC ATAGGGGAGC CCTTATTGCA ACCTCAGCAG 1860
 TTCTCTTTTT CCTTATCTC TTGATTAATG CCTACTTTGC CTAATTGAAA GGAAAATCAT 1920
 CTTATGAGTA AATACCTGCT AAAACTTCTC GTTTATGTGTT TTTCAGCTTT AACCTTTGGC 1980
 TCTCTCTTTT TAATCATTTG TTTTATCCTC ATCAAAGGCT TACCTCATCT AAGTCTATCC 2040

CTCTTTTCTT	GGACTTATAC	TTCTGAGAAC	ATTTCCCTTA	TGCCAGCGAT	TATTTCCACC	2100
GTTATTTCTGG	TCTTTGGTGC	TCTTCTTTTA	GCCTTGCCCA	TAGGGATTTT	TGCTGGTTTT	2160
TATCTTGTGG	AATATACAAA	AAAAGATPCC	CTTTGTGTTA	AAATCATGCG	ATTGGCCTCA	2220
GATACCTTAT	CTGGGATTCC	TTCCATTGTT	TTTGGTCTGT	TTGGCATGCT	CTTCTTTGTA	2280
GTCTTCTTAG	GTTTTCAATA	CTCTCTGTTA	TCAGGAATCT	TAACCTCAGT	TATCATGGTG	2340
TTGCCAGTCA	TTATTCGCTC	AACAGAAGAA	GCCCTTTTAT	CTGTTAGTGA	TAGCATGCGT	2400
CAAGCAAGTT	ATGGACTTGG	GGCAGGTAAG	TTACGGACTG	TTTTTAGAAT	TGTTCTACCA	2460
GTTGCCATGC	CAGGTATTTT	AGCTGGAGTG	ATACTAGCTA	TTGGCCGTAT	CGTTGGTGAA	2520
ACAGCTGCCC	TCATGTATAC	ATTAGGTACC	TCTACCAATA	CGCCAAGTAG	TCTCATGTCT	2580
TCAGGCCGTT	CTCTAGCCCT	ACATATGTAT	ATGCTGTCAA	GTGAGGGGCT	ACATGTCAAT	2640
GAAGCCTATG	CTACCGCGT	GATTTTGATT	ATTACTGTTT	TAATGATAAA	TACTCTATCA	2700
AGCTTATTAT	CTCGAAAAC	TGTGAAAGGA	GCTTCCTAGT	ATGGGAACAT	TTTCAGTCAG	2760
ACACCTAGAC	TTATTTTACG	GGGATTTTCA	AGCCTTAAAA	AATATTTTCA	TTCAATTACC	2820
AGAAAGACAG	ATTACTGCCT	TGATAGGCCC	ATCTGGTTGT	GGCAAATCAA	CTTTTCTAAA	2880
AACCCCTAAC	CGGATGAACG	ATTTGGTTCC	TTCTTGCCAT	ATTGAAGGCC	AAGTCTCTT	2940
AGATGAGCAA	GATATTTATA	GTAGCAAATT	CAACCTTAAT	CAGCTACGTA	AGCGTGTAGG	3000
GATGGTTTTT	CAACAGCCTA	ATCCCTTTGC	CATGTCTATC	TATGATAACG	TGGCTTATGG	3060
CCCAAGGACA	CATGGTATTC	GAGACAAAA	ACAATTAGAT	GCCTTAGTGG	AGAAATCTTT	3120
AAAAGGGGCA	GCCATTTGGG	AAGAAGTCAA	AGATGATCTT	AAAAAGAGTG	CCATGTCCTT	3180
ATCTGGCGGT	CAGCAGCAAC	GCCTTTGCAT	TGCGCGAGCT	TTAGCAGTAG	AACCTGATAT	3240
TCTGTTAATG	GATGAGCCGA	CTTCAGCCTT	AGACCCATC	TCCACTTTAA	AAATTGAAGA	3300
CCTCATTCAG	CAACTAAAA	AGGATTATAC	GATTATCATT	GTTACCCATA	ACATGCAACA	3360
AGCTTCACGT	ATTTTCAGATA	AAACTGCTTT	TTTCTTAACA	GGAGAAATTT	GCGAATTTGG	3420
AGATACCGTT	GACGTGTTTA	CCAATCCAAA	AGATCAGCGC	ACAGAAGACT	ATATTTCAGG	3480
ACGGTTCGGA	TAAGGAAGGA	AAAACCTATG	AGAAATCAAT	TTGACTTAGA	ATTGCATGAA	3540
TTAGAACAAT	CCTTTTTAGG	ACTAGGGCAA	CTTGTCCTTG	AAACAGCTTC	AAAAGCCTTA	3600
CTGGCCTTAG	CCTCCAAAGA	CAAGGAGATG	GCAGAGCTAA	TTATCAATAA	GGATCATGCT	3660
ATCAACCAAG	GTCAAAGCGC	TATCGAATTG	ACCTGTGCC	GTTTGTGGC	CTTGCAAGCAG	3720
CCACAAGTGT	CTGACCTTCG	ATTTGTGATT	AGCATCATGT	CTTCTTGTTT	AGACCTTGAA	3780

CGTATGGGAG	ACCATATGGC	AGGCATTGCC	AAAGCTGTTT	TGCAACTAAA	AGAAAATCAA	3840
CTAGCCCCTG	ACGAAGAACA	GTTACACCAA	ATGGGTAAAT	TATCCCTCAG	CATGCTAGCC	3900
GATTTATTGG	TTGCCTTTCC	TTTGCACCAA	GCCTCAAAAG	CTATTAGTAT	TGCTCAAAAA	3960
GATGAACAGA	TTGACCAATA	TTATTATGCC	TTATCAAAGG	AAATCATTGG	ACTTATGAAA	4020
GACCAAGAAA	CCTCAATFCC	CAATGGAACT	CAATACCTTT	ATATCATAGG	GCATCTGGAA	4080
CGCTCGCTGA	TTACATGCT	AACATTTGTG	AACGCCTAGT	CTACCTAGAA	ACAGGAGAAC	4140
TAGTGGATTT	GAATTAATTC	AACATAATCCT	TAAAAGAGAA	GAGTACGATT	AAGTACTCTT	4200
TTTTATGGTT	GTA AAAAAGT	TCATTTGACC	AATTTAAGCA	GTGTAGATAG	TGAGGAGTTG	4260
TTTCAATFCT	ATCGTGAACG	AGGGAATGCT	GAAAACFTTA	TCAAAGAAAAG	GAAAGCAGGA	4320
TTCTTTGGGG	ATAAGACAGA	TAGTTCGACC	ATGATTAAGA	ATGAAGTACG	TATGATGATG	4380
GGCTGTCTGG	CTTATAATCT	CTACCTCTTT	TTAAAGCAGC	TAGCTGGTGA	TGAAGTAAAG	4440
TCCTTGACTA	TCAAGCGTTT	TCGACGTCTC	FTCCTTCATA	TTGCCGGAAA	ATATGTCTCT	4500
ACTGCTAGAC	GACATAFTCT	CAAATFCTCA	AGTCTATACG	CCTATFCAA	ACAGTTTCAA	4560
GCCTTATTTG	ATACAATCTG	CCAGATAAAAT	CTGATACTCC	CTGTTCCATA	TAGAGCTAGA	4620
GGGCAGGGGA	AAACATGCCT	AACAGAATAA	GTCACCTTAT	TTTAAAAATC	GAGCATCAAA	4680
CCAAGGGAGG	AGTCTGCCCT	TTTTTAGGAA	AAAATCAAGA	CAAATCTCCT	CAATFATGTC	4740
TCGAACATCA	GAAATFAAGC	AAAATCACCA	GAAGGACAGT	ATTTCAACTA	GCTTTTCTGG	4800
TAATTTTGA	ACTGTGAGT	TCGTTAGTGC	CAGATATGAA	TAATTTGGGA	TGATAAATCT	4860
TTCTTCCTCA	GGTAGCCTAT	CATAATACTC	FTCAAAAATC	TTATCAAAAA	CACTCTCTTT	4920
CTTTTGGGGC	ATAGTTTCAT	CTTCGTATGT	AGGAGTCCTC	ATCAAGAAAT	ACTTCAATTC	4980
TAGGTATFCC	TTATCCAACT	CTATATAACT	TGGCATCAAC	TTGTAATCTT	CAACCCCAA	5040
ACGFTCAGCA	ATATATTTTA	ACTTTGTTAG	TATTGGTCTG	GATFCTCCAT	FTTCAATFCT	5100
AATTAATFGA	CGGATACTTA	ATTCAGACTC	ATCACCACAA	AATFCTGAAC	GACTGATFI	5160
TTTAGCCAAA	CGTAATCTTT	TAATTTTFTC	GCCAAACTCT	CGCAACCTAC	AAGAACTFCC	5220
TGAGTTGTTT	ACCTCTATTA	TAAGCATATA	CTGAATCAAA	CTATCTATCA	GATTTCTTCT	5280
CACTTFAACT	AAAGACTAAG	AGTTTATCCC	FTCGTCTCGG	TTTTTGTGTA	TTTTTCCACC	5340
ATACCCAGT	AATGCAAGTG	CAAAATCCCC	TAGAATATGA	TAGAATAAGA	GAAAGAACTC	5400
TATCAAGGAG	GAAATCATGG	AAAAACAAAC	CGTCGCCGTC	TTGGGGCCTG	GTTCTTGGGG	5460
AACCGCCCTT	TCACAAGTCT	TAAATGACAA	TGGACACGAG	GTACGTATTT	GGGGAAATCT	5520
TCCCAGACAA	ATCAATGAAA	TTAATACACA	CCATACTAAT	AAGCACTACT	TTAAAGATGT	5580

995

CGTTCTAGAC	GAAAATATCA	TTGCCTACAC	CGACTTAGCA	GAAACATTGA	AAGATGTGGA	5640
TGCGATTTTG	TTTGTGTGCC	CAACAAAAGT	GACACGACTT	GTTGCCCAGC	AAGTTGCACA	5700
AACCTTGGAC	CATAAGGTTA	TCATCATGCA	CGCATCAAAG	GGATTAGAAC	CTGATAGCCA	5760
TAAACGATTA	TCAACCATTC	TTGAAGAAGA	AATTCCTGAA	CATCTCCGTA	GTGATATCGT	5820
CGTTGTTTCA	GGGCCTAGTC	ATGCAGAAGA	GACCATTGTG	CGTGACCTAA	CTTTAATAAC	5880
TGCTGCTTCT	AAAGATTTAC	AAACAGCTCA	ATACGTTTAC	AAGCTATTTA	GTAATCACTA	5940
CTTCCGACTT	TATACCAATA	CGGATGTTAT	CGGGGTTGAA	ACTGCTGGTG	CTCTTAAAAA	6000
TATTATGCT	GTCGGTGCTG	GAGCTTTACA	TGCTTGTGGA	TTTGGTGATA	ATGCTAAGGC	6060
AGCCATCATC	GCTCGAGGTT	TAGCAGAAAT	CACCCGCCTA	GGGGTAGCAC	TCGGGGCCAG	6120
TCCATTGACC	TATAGCGGCT	TATCTGGTGT	GGGAGATTTG	ATCGTAACGG	GAACTTCCAT	6180
CCACTCTCGT	AACTGGAGAG	CTGGAGATGC	TCTCGGACGA	GGAGAATCCC	TAGCTGATAT	6240
AGAAGCTAAT	ATGGGCATGG	TAATCGAAGG	AATTTCAACG	ACTCGAGCAG	CCTATGAACT	6300
AGCCCAAGAA	CTTGGAGTCT	ATATGCCCAT	TACACAGGCT	ATTTACCAAG	TTATTTATCA	6360
CGGAACCAAT	ATCAAAGATG	CCATTTATGA	CATCATGAAC	AATGAATTTA	AAGCAGAAAA	6420
TGAGTGGTCT	TAACCCCTTA	TAGAAAGGAT	TTTTATGACA	TCAAAGTTA	GAAAGGCAGT	6480
CATCCCTGCT	GCTGGACTAG	GAACTCGATT	TTTACCAGCA	ACCAAGGCC	TTGCCAAAGA	6540
AATGTTGCCA	ATCGTAGACA	AACCAACTAT	CCAGTTTATC	GTGGAAGAAG	CTCTCAAATC	6600
AGGTATTGAA	GATATTCTAG	TTGTCACTGG	TAAATCAAAA	CGTTCTATTG	AGGACCACTT	6660
TGATTCAAAC	TTCGAATTGG	AATATAACCT	CAAAGAAAAA	GGGAAAACAG	ATCTTTTGAA	6720
GCTAGTTGAT	AAAACAACCTG	ACATGCGTCT	GCATTTTATC	CGCCAAACTC	ATCCACGCGG	6780
TCTCGGAGAT	GCTGTTTTGC	AAGCCAAGGC	TTTCGTCCGA	AATGAACCTT	TTGTCTGTTAT	6840
GCTTGGTGAT	GACTTGATGG	ATATCACAGA	CGAAAAGGCT	GTTCCACTTA	CCAAACAACCT	6900
CATGGATGAC	TACGAGCGTA	CCCACGCGTC	TACTATCGCT	GTCATGCCAG	TCCCTCATGA	6960
CGAAGTATCT	GCTTACGGGG	TTATTGCTCC	GCAAGGCGAA	GGAAAAGATG	GTCTTTACAG	7020
TGTTGAAACC	TTTGTTGAAA	AACCAGCTCC	AGAGGACGCT	CCTAGCGACC	TTGCTATATAT	7080
CGGACGCTAC	CTCCTCACGC	CTGAAATTTT	TGAGATTCTC	GAAAAGCAAG	CTCCAGGTGC	7140
AGGAAATGAA	ATTCAGCTGA	CAGATGCAAT	CGACACCCTC	AATAAACAC	AACGTGTATT	7200
TGCTCGTGAG	TTCAAAGGGG	CTCGTTACGA	TGTCGGAGAC	AAGTTTGGCT	TCATGAAAAC	7260
ATCCATCGAC	TACGCCCTCA	AACACCCACA	AGTCAAAGAT	GATTTGAAGA	ATTACCTCAT	7320

996

CCAACCTTGA	AAAGAATTGA	CTGAGAAGGA	ATAACAAAAT	CATTTATATA	AAGATTAGCC	7380
ACACATAAAT	TAAGTAAATT	CTCTACTTGA	ATCTACCTAT	TTAATAAAAA	CTAATGAAAA	7440
CGCTATACTT	GTATTTGTTT	TTTCATTAAA	ATAAGAGTAG	AATAAAATAG	TATAGTAAAA	7500
CAAAAAAGCA	CCGAATCGGT	GCGCACTTTT	TCAAGTTGTG	TACGGACAAA	GCCTTATTTT	7560
AACTTTGCTA	TGTTGTTTCT	AATGGTTCCA	AAATAATAAA	TAATTTTAAA	TTTGACTTAA	7620
CTGTTGGAGT	AGTCATGGTT	AAATTAATC	AACCGAGCCG	AACATAAGTT	GTTTAATTTT	7680
GTGGAAGCTA	TTAATAAAAA	TATAATAAGG	GAGAAAGATA	GGTGAATTT	TAATTTTAAA	7740
GTAATTGCGG	ACACTATCAA	AGAAAAAGAT	TATGGAGAAC	AAATTTGTAG	AATTTATCGA	7800
AAACAATAAA	AAAGTAATCA	TTTCATCAGT	TGCAGTTGGT	GTTGTATTGG	TATTAGGGTT	7860
TGGATGGTAT	TCATATAACC	AACAACAAGC	AGAACAACAA	GCAAAAATTG	TACAATTAGA	7920
AAAAGATAGC	AAATCAGACA	AAGAACAAGT	TGATAAACTA	TTTGAATCAT	TTGATGCATC	7980
TTCAGATGAA	TCTATTTCTA	AATTAAGA	ACTATCTGAA	ACTTCACTTA	AAACCGATGC	8040
AGGTAAAGAC	TATCTTAATA	ACAAAGTCAA	AGAATCATCT	AAAGCAATTG	TAGATTTTCA	8100
TTTGCAAAAA	GGTTTGCTT	ATGATGTAA	AGATTCAGAT	GACAAATTTA	AAGATAAAGC	8160
AACTCTTGAA	ACAAATGTAA	AAGAAATTAC	AAAACAAATT	GATTTTATCA	AAAAAGTTGA	8220
TGAAACTTTT	AAACAAGAGA	ATTTGGAAGA	AACTCTTAAA	TCTCTAAATG	ATCTTGTTGA	8280
TAAATATCAA	AAACAAATCG	AACTTTTGAA	GAAAGAAGAA	GAAAAAGCTG	CTGAAAAAGC	8340
TGCTGAAAAA	GCAAAGGAAT	CTTCTAGTCA	AAGTAATCT	TCTGGTAGTG	CTTCTAATGA	8400
GTCTTATAAT	GGATCTTCCA	ATTCAAATGT	AGATTATAGT	TCATCTGAAC	AAACTAATGG	8460
ATATTCAAAT	AATTATGGCG	GTCAAGATTA	TTCTGGTTCA	GGAGATAGTT	CAACAAATGG	8520
TGGATCATCA	GAACAATATT	CATCTAGCAA	TTCAAACAGC	GGAGCAAATA	ATGTCTACAG	8580
ATATAAAGGC	ACTGGTGCTG	ACGGCTATCA	AAGATACTAC	TACAAAGATC	ATAATAATGG	8640
AGATGTGTAT	GATGACGATG	GAAATTACCT	TGGGAACTTT	GGTGGCGGCA	TTGCAGAACC	8700
TAGTCAACGC	TAATAACTAT	TTTAGAGCTG	TGTTGTTTCG	AATGGTTCCA	AAACACATTA	8760
AAAGCTACTC	ATTTTTTAAG	TAGCTTTTTT	CTTATCAAG	TTTACATATT	ATACTCAATG	8820
AAAATCAAAT	TCAAACCACG	TCAGCATCGC	CTTACCGTAG	GTATGGTTAC	TGACTTCGTC	8880
AGTTTCATCT	ACAACCTCAA	AACCATGTTT	TGAGCTGACT	TCGTCAGTTC	TATCTACAAC	8940
CTCAAAGCAG	TGCTTTGAGC	AACCTGCGGC	TAGCTTCCTA	GTTTGCTCTT	TGATTTTCAT	9000
TGAGTATTAG	TCGTCACAAT	CCCATCCCT	TGTAGAAAAG	CAAAATGGCG	AGTCCTACGA	9060
ACAAGACTAC	CGCTCCTAAT	CTCTGGCTGG	TGTTATACAT	CCGTTTTTCT	CCTCTAACTG	9120

GAAAGATAAC TGCTAGAAAT GCGCCACCAA CTGCACCACC GATATGGCCT GCTAGGCTGA	9180
TTCTTGGAAT CAGAACACTT CCAATAATGT TAACCACAAA AAGTGCAGA TAGGATTGCC	9240
CTAGCTGTTG GATATAAGGA TTGCGAGTTG CATAGCGAAG AACATAATC GCGGCAAATA	9300
GCCCATAAAG AGAGGTAGAG GCGCCTGCTG CTAAGGATTT AGGACTAAAT AAAAAACAA	9360
AGAGATTGCC CATCATTCCT GATAAAAGAT AGAGAAAGAA AAAC TGCTTA GAACCGAAAA	9420
TCTCCTCTAC CTGCCTTCCA AGATAATAAA GTGAAAGCAT ATTAACAATG AAATGTTCCC	9480
ACCCAATATG AACAAAAATG GCAGACAAGA GACGCCAAC CTGCTCGGGA AAGAGGCGAA	9540
TAGCTGGCCC ATACATGGCT CCAAATCGAA ATAATGTATC TGCCCTGTCA AAGTTCCGC	9600
CTGCAGTGAC CAACATTAGT AAAAATACCA AGGCCGTAC TAAGAGGAAG AAAC TCGTCA	9660
CAGGGTAACG TCTATCAAAG ATTTCTTCA TCAATTAATA CCTCCTGAAC AGGAATATCA	9720
TGGTTTTCAG GTATAAAGTC CTGAATTTGA CAAGGATATA TCGTACTCAA AGTACGACCA	9780
GAAAAATGTT CCAGATAGCG GTCATAATAG CCTCCACCGT ATCCTATCCG ATATCCTTTC	9840
GTCTGAAAAG CCAGACCAGG AACATGAATC AAATCAATCT GAGATGCATC CACCACTTCC	9900
AAATCTCCCT GTAGCTCCAG TAAGGCAAAG AAAGTTTTTA CCAACTGTTG CGGATCATAG	9960
ACCACAAAGT CCATGCGCCC CTGCGGATAA GTTTTGGGTA TTAAAACCTT CTGCCGTCC	10020
TTCAGCGCCT GCTCAATCAG TTCCTGCGTT TGAAACTCAT GAGAAAAAGA GAGGTAGGTT	10080
GCGATGACCT TGGCTTCTTG ATAAAAGGGG TGTGTAAAA GCCGCTCGGT TAAAGCTTGG	10140
TCTATAGCCT GTTTTTGCTC TTGAGATATA GCCTTCATTT CATGCAAGAC TTGCTTGCCT	10200
AAATCCGATT TCATAGACAA GCCCTCTATT CTGCTGCCCTT CTTTTTCAGG AACTAGACA	10260
CCGCAGCCAC CCAATAGCT AAGACTTCTT CCTTAGGACT CATTTGAGGG TGATGAAGAG	10320
CGTAGGGACT ATCGATACCT AGCCAAAACA TCACGCCATC AACCTTTGAA AGGAGATAAC	10380
CAAAGTCCTC GCCTGTCATA GCAGGTTCGA TATCAATCAA CTCGATTCCG TCTTTTTCGT	10440
CAAAGAAGTC CATCAGTTCA CGCGCCAAGG CTGGATTGTT CTCAACAGGT AGGTATCCAC	10500
CTTGTTTGAG TTCCACTTCG ACTTCCATAT CAAAGGCAGC TGCAACCCCT TCTGCAACTG	10560
TTTTTACCCT CTTTTGCACC AAGAGACTCA TGTCCTGTGT CAAGGCACGA ATAGTTCCAT	10620
GTAAAAAAGC TGTGTCTGTG ATGACATTGT TGGTGGTTCC AGCTTGAAAA ACGCCGAAGG	10680
TCACCACTGC TCCCTCGATT GGGTTGACAT TGCGGCTAAC AACTGACTGC ACTTGGGTCA	10740
CAAAGTAACT AGCCGCCACC AAGGCGTCAT TGGCTTCATG AGGAAAAGCT GCGTGGCCAC	10800
CTTTGCCTTT GAAACGGATC TTCACCTCGC AAGTTCCTGC AAAGAGTGTA TGAGTATTAG	10860

998

TCGCAATCTG GCCGACTTTC AAATCTGGAC GAACATGGAG ACCATAGAAT TGATCTGGCA 10920
 ACCAATCTCC AAAAGCACCG TCCTCATACA TGAGCATACC ACCAGCTTCA TTTTCTTCAG 10980
 CAGGCTGAAA TAGAAAGAGC AGATTATTCT TGGGTGTGTC CTCAAGGGCG CGCTCAAGAC 11040
 AGCCTAAGGC AATGGTCATA TGAAAATCAT GGACACAGGC ATGCATGCGA CCTTGGTGTT 11100
 GAGAAGCAAA AGGTAGACCT GTTTGTTCGA CGATAGGCAG GCCATCAATA TCTGTCCGCC 11160
 AACCAATGGT TCGCTCCGGC TGACTTCCCT GCAGGTAGAC CAAAATCCCT GTCCGCCAAG 11220
 TACGAATTTG AACAAAATCC TTGCCCGTAG TCAATTTCTC AATCACATCC AGCAAATAAG 11280
 CCTGAGTCTT GAACTCCTCC AAGCCAATCT CTGGAATCTG GTGTAAATCT CGTCTAGTCT 11340
 GAATCAAATC TAACATCTAT CTGTCTCCG ATATAGCAGA AAGAGGCTGG AAAAAGGGTT 11400
 CCGCTCTTTT TTTACTTTTA CAATTACAAG GTACGAAGCG CATCCTCTAG CGCTGTTTTT 11460
 TGTGAGTTT GGGCATCAAT TTCTTTGATA ATACGAGCTG GAACACCTGC TACTACCACG 11520
 TTTTCTGGGA CATCTGGGT AACAAATAGCT CCTGCTGCGA CAACTGAACC ACTACCGATT 11580
 TGGACTCCTT CGATAACCAC TGCATTAGCA CCGATAAGAA CATTGTCTCC GACACGGACT 11640
 GGTTCAGCAC TAGCTGGCTC AATCACACCT GCCAAAATG CACCTGCACC AACGTGGCTA 11700
 TTTTTTCCAA CGATGGCAGC GCCACCAAGG ATGGCACCCA TGTC AATCAT GGTTCAGCA 11760
 CCGATTTTCCAG CACCGATATT GATAACAGAT CCCATCATGA TAACAGCATT GTCACCAATT 11820
 TCCACCTGGT CACGGATAAT CGCACCTGGC TCGATACGAG CGTTGATAGC ACGCTTATCT 11880
 AGCAAAGGAA CTGCAGAATT ACGAGCATCT TGCTCGACAA CATAATCTTG ATTTTCTACC 11940
 AAACCTTCAA GAAGCGGAGC CACATCCTTC CAGTCTCCGA ATAGGACATT TCCTAGTTTG 12000
 ACAACAGAGC TAGGCACAGC AGTTGCGAGT TGCCCTCAA AGTTACTTTT GACTGTTGTT 12060
 TTCTTTTCAG CATTGGCGAT AAATTGGATA ATTTCTTGAG CGTTCATTTT TGTAGCAGTC 12120
 ATAGGTG 12127

(2) INFORMATION FOR SEQ ID NO: 149:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 12566 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 149:

CCATCCTTCT GTTGATGTGA CAGGAATGAT GATAAATCAA CCAGTAGCTA GTCGCGAAGA 60
 GGTGACAGAG GCTTTGAGTC ACTTGGCGGT AGAGACAAT AGTCTCATTG CTCGTCGAAT 120